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Grimaldi, Christopher Gurney, Austin L. Hillan, Kenneth, J. Kljavin, Ivar J. Mather, Jennie P. Pan, James Paoni, Nicholas F. Roy, Margaret Ann Stewart, Timothy A.

JAN 2 5 2001

Tumas, Daniel
Williams, P. Mickey
Wood, William, I.

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Pro Thr Leu Gly Val Val Pro Gln Ala Ser Val Pro Leu Leu Thr
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Asp Leu Ala Gln Trp Glu Pro Val Leu Val Pro Glu Ala His Pro 80 85 90

Asn Ala Ser Leu Thr Met Tyr Val Cys Thr Pro Val Pro His Pro 95 100 105

Asp Pro Pro Met Ala Leu Ser Arg Thr Pro Thr Arg Gln Ile Ser 110 115 120

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Cys Cys Cys Phe His Gly Pro Ala Phe Ser Thr Leu Asn Pro Val

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Ile Ile Asp Ser Ser Arg Ser Val Asn Thr His Asp Tyr Ala Lys
65 70 75

Val Lys Glu Phe Ile Val Asp Ile Leu Gln Phe Leu Asp Ile Gly
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Pro Asp Val Thr Arg Val Gly Leu Leu Gln Tyr Gly Ser Thr Val 95 100 105

Lys Asn Glu Phe Ser Leu Lys Thr Phe Lys Arg Lys Ser Glu Val 110 115 120

Glu Arg Ala Val Lys Arg Met Arg His Leu Ser Thr Gly Thr Met 125 130 135

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Phe	Glu	Gly	Tyr	Ile 515	Leu	Arg	Glu	Asp	Gly 520	Lys	Thr	Cys	Arg	Arg 525
Lys	Asp	Val	Cys	Gln 530	Ala	Ile	Asp	His	Gly 535	Cys	Glu	His	Ile	Cys 540
Val	Asn	Ser	Asp	Asp 545	Ser	Tyr	Thr	Cys	Glu 550	Cys	Leu	Glu	Gly	Phe 555
Arg	Leu	Ala	Glu	Asp 560	Gly	Lys	Arg	Cys	Arg 565	Arg	Lys	Asp	Val	Cys 570
Lys	Ser	Thr	His	His 575	Gly	Cys	Glu	His	Ile 580	Cys	Val	Asn	Asn	Gly 585
Asn	Ser	Tyr	Ile	Cys 590	Lys	Cys	Ser	Glu	Gly 595	Phe	Val	Leu	Ala	Glu 600
Asp	Gly	Arg	Arg	Cys 605	Lys	Lys	Cys	Thr	Glu 610	Gly	Pro	Ile	Asp	Leu 615
			Arg	605	_	_			610					615
Val	Phe	Val		605 Asp 620	Gly	Ser	Lys	Ser	610 Leu 625	Gly	Glu	Glu	Asn	615 Phe 630
Val	Phe Val	Val Val	Ile	Asp 620 Gln 635	Gly Phe	Ser Val	Lys Thr	Ser Gly	610 Leu 625 Ile 640	Gly	Glu Asp	Glu Ser	Asn Leu	Phe 630 Thr 645
Val Glu Ile	Phe Val Ser	Val Val Pro	Ile Lys	605 Asp 620 Gln 635 Ala 650	Gly Phe Ala	Ser Val Arg	Lys Thr	Ser Gly Gly	610 Leu 625 Ile 640 Leu 655	Gly Ile Leu	Glu Asp Gln	Glu Ser Tyr	Asn Leu Ser	Phe 630 Thr 645 Thr 660
Val Glu Ile	Phe Val Ser	Val Val Pro	Ile Lys Lys	605 Asp 620 Gln 635 Ala 650 Glu 665	Gly Phe Ala Phe	Ser Val Arg	Lys Thr Val	Ser Gly Gly Arg	610 Leu 625 Ile 640 Leu 655 Asn 670	Gly Ile Leu Phe	Glu Asp Gln Asn	Glu Ser Tyr Ser	Asn Leu Ser	Phe 630 Thr 645 Thr 660 Lys 675
Val Glu Ile Gln Asp	Phe Val Ser Val	Val Val Pro His	Ile Lys Lys Thr	Asp 620 Gln 635 Ala 650 Glu 665 Ala 680	Gly Phe Ala Phe Val	Ser Val Arg Thr	Lys Thr Val Leu	Ser Gly Gly Arg	610 Leu 625 Ile 640 Leu 655 Asn 670 Lys 685	Gly Ile Leu Phe	Glu Asp Gln Asn Met	Glu Ser Tyr Ser	Asn Leu Ser Ala	Phe 630 Thr 645 Thr 660 Lys 675 Gly
Val Glu Ile Gln Asp	Phe Val Ser Val Met	Val Val Pro His Lys	Ile Lys Lys Thr	Asp 620 Gln 635 Ala 650 Glu 665 Ala 680 Leu 695	Gly Phe Ala Phe Val	Ser Val Arg Thr Ala	Lys Thr Val Leu His	Ser Gly Gly Arg Met	610 Leu 625 Ile 640 Leu 655 Asn 670 Lys 685 Met 700	Gly Ile Leu Phe Tyr	Glu Asp Gln Asn Met	Glu Ser Tyr Ser Gly	Asn Leu Ser Ala Lys Ser	Phe 630 Thr 645 Thr 660 Lys 675 Gly 690 Phe 705
Val Glu Ile Gln Asp Ser	Phe Val Ser Val Met Met	Val Val Pro His Lys Thr	Ile Lys Lys Thr Lys	Asp 620 Gln 635 Ala 650 Glu 665 Ala 680 Leu 695 Gly 710	Gly Phe Ala Phe Val Ala Ala	Ser Val Arg Thr Ala Leu Arg	Lys Thr Val Leu His Lys	Ser Gly Gly Arg Met His	610 Leu 625 Ile 640 Leu 655 Asn 670 Lys 685 Met 700 Ser 715	Gly Ile Leu Phe Tyr Phe	Glu Asp Gln Asn Met Glu Arg	Glu Ser Tyr Ser Gly Arg	Asn Leu Ser Ala Lys Ser Pro	Phe 630 Thr 645 Thr 660 Lys 675 Gly 690 Phe 705 Arg 720

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 Leu Pro Lys Thr Val Gln Gln Pro Thr Glu Ser Glu Pro Val Thr
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 His Arg Tyr Leu Phe Glu Glu Asp Asn Leu Leu Arg Ser Thr Gln
 Lys Leu Ser His Ser Thr Lys Pro Ser Gly Ser Pro Leu Glu Glu
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 Lys His Asp Gln Cys Lys Cys Glu Asn Leu Ile Met Phe Gln Asn
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35 40 45

Gly Glu Val Val Leu Pro Ala Trp Tyr Thr Leu His Gly Glu Val
50 55 60

Ser Ser Ser Gln Pro Trp Glu Val Pro Phe Val Met Trp Phe Phe
65 70 75

Lys Gln Lys Glu Lys Glu Asp Gln Val Leu Ser Tyr Ile Asn Gly 80 85 90

Val Thr Thr Ser Lys Pro Gly Val Ser Leu Val Tyr Ser Met Pro 95 100 105

Ser Arg Asn Leu Ser Leu Arg Leu Glu Gly Leu Gln Glu Lys Asp 110 115 120

Ser Gly Pro Tyr Ser Cys Ser Val Asn Val Gln Asp Lys Gln Gly

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Pro	Ala	Val	Gln		Gln	Trp	Asp	Arg		Leu	Pro	Ser	Phe	
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Leu	Thr	Asn	Leu	Ser 215	Ser	Ser	Met	Ala	Gly 220	Val	Tyr	Val	Cys	Lys 225
Ala	His	Asn	Glu	Val 230	Gly	Thr	Ala	Gln	Cys 235	Asn	Val	Thr	Leu	Glu 240
Val	Ser	Thr	Gly	Pro 245	Gly	Ala	Ala	Val	Val 250	Ala	Gly	Ala	Val	Val 255
Gly	Thr	Leu	Val	Gly 260	Leu	Gly	Leu	Leu	Ala 265	Gly	Leu	Val	Leu	Leu 270
Tyr	His	Arg	Arg	Gly 275	Lys	Ala	Leu	Glu	Glu 280	Pro	Ala	Asn	Asp	Ile 285
Lys	Glu	Asp	Ala	Ile 290	Ala	Pro	Arg	Thr	Leu 295	Pro	Trp	Pro	Lys	Ser 300
Ser	Asp	Thr	Ile	Ser 305	Lys	Asn	Gly	Thr	Leu 310	Ser	Ser	Val	Thr	Ser 315
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Leu	Thr	Pro	Thr	Pro 335	Ser	Leu	Ser	Ser	Gln 340	Ala	Leu	Pro	Ser	Pro 345
Arg	Leu	Pro	Thr	Thr 350	Asp	Gly	Ala		Pro 355	Gln	Pro	Ile	Ser	Pro 360
Ile	Pro	Gly	Gly	Val 365	Ser	Ser	Ser	Gly	Leu 370	Ser	Arg	Met	Gly	Ala 375
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Met Gly Phe Ser Gly Asn Gly Val Thr Ile Cys Glu Asp Asp Asn 50 55

Glu Cys Gly Asn Leu Thr Gln Ser Cys Gly Glu Asn Ala Asn Cys
65 70 75

Thr Asn Thr Glu Gly Ser Tyr Tyr Cys Met Cys Val Pro Gly Phe 80 85 90

Arg Ser Ser Ser Asn Gln Asp Arg Phe Ile Thr Asn Asp Gly Thr 95 . 100 . 105

Val Cys Ile Glu Asn Val Asn Ala Asn Cys His Leu Asp Asn Val
110 115 120

Cys Ile Ala Ala Asn Ile Asn Lys Thr Leu Thr Lys Ile Arg Ser 125 130 135

Ile Lys Glu Pro Val Ala Leu Leu Gln Glu Val Tyr Arg Asn Ser 140 145 150

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Ser	Ala	Lys	Asp	Thr 185	Leu	Ser	Asn	Ser	Thr 190	Leu	Thr	Glu	Phe	Val 195
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Asp	Lys	Leu	Ser	Val 215	Asn	His	Arg	Arg	Thr 220	His	Leu	Thr	Lys	Leu 225
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Gln	Lys	Thr	Thr	Glu 245	Phe	Asp	Thr	Asn	Ser 250	Thr	Asp	Ile	Ala	Leu 255
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Вси	Leu	гуs		320	ASII	*1*	Asp	ASII	325				Olu	330
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Val Thr Lys Ser Leu	Ile Leu Val Pro	Ser Tyr Thr Asp	Ser Glu Asp	320 Val 335 Leu 350 Arg 365 Met 380 Asn 395	Ile Glu Tyr Asn Glu	Ser Lys Arg Gly	Val Ile Ser Ser	Ser Thr Leu Trp	325 Met 340 Phe 355 Cys 370 Ser 385 Ser 400	Ser Thr Ala Ser Cys	Ser Leu Phe Glu	Asn Ser Trp Gly	Pro His Asn Cys	330 Pro 345 Arg 360 Tyr 375 Glu 390 His 405
Val Thr Lys Ser Leu Leu	Ile Leu Val Pro Thr	Ser Tyr Thr Asp Tyr	Ser Glu Asp Thr	320 Val 335 Leu 350 Arg 365 Met 380 Asn 395 Ala 410	Ile Glu Tyr Asn Glu Ile	Ser Lys Arg Gly Thr	Val Ile Ser Ser His	Ser Thr Leu Trp Thr	325 Met 340 Phe 355 Cys 370 Ser 400 Ser 415	Ser Thr Ala Ser Cys	Ser Leu Phe Glu Arg	Asn Ser Trp Gly Cys Ser	Pro His Asn Cys Asn Ile	330 Pro 345 Arg 360 Tyr 375 Glu 390 His 405 Gly 420

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 Ile Asn Thr Asn Thr Asn Lys Leu Phe Cys Ser Ile Ile Ala Gly
                 485
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                 500
 Glu Gly Ile His Leu Tyr Leu Ile Val Val Gly Val Ile Tyr Asn
 Lys Gly Phe Leu His Lys Asn Phe Tyr Ile Phe Gly Tyr Leu Ser
 Pro Ala Val Val Gly Phe Ser Ala Ala Leu Gly Tyr Arg Tyr
 Tyr Gly Thr Thr Lys Val Cys Trp Leu Ser Thr Glu Asn Asn Phe
 Ile Trp Ser Phe Ile Gly Pro Ala Cys Leu Ile Ile Leu Val Asn
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 Ala Gly Leu Lys Pro Glu Val Ser Cys Phe Glu Asn Ile Arg Ser
 Cys Ala Arg Gly Ala Leu Ala Leu Leu Phe Leu Leu Gly Thr Thr
 Trp Ile Phe Gly Val Leu His Val Val His Ala Ser Val Val Thr
 Ala Tyr Leu Phe Thr Val Ser Asn Ala Phe Gln Gly Met Phe Ile
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                                      655
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Gly Pro His Val His Tyr Gly Trp Gly Asp Pro Ile Arg Leu Arg
35 40 45

His Leu Tyr Thr Ser Gly Pro His Gly Leu Ser Ser Cys Phe Leu 50 55 60

Arg Ile Arg Ala Asp Gly Val Val Asp Cys Ala Arg Gly Gln Ser
65 70 75

Ala His Ser Leu Leu Glu Ile Lys Ala Val Ala Leu Arg Thr Val 80 85 90

Ala Ile Lys Gly Val His Ser Val Arg Tyr Leu Cys Met Gly Ala 95 100 105

Asp Gly Lys Met Gln Gly Leu Leu Gln Tyr Ser Glu Glu Asp Cys 110 115 120

Ala Phe Glu Glu Ile Arg Pro Asp Gly Tyr Asn Val Tyr Arg
125 130 135

Ser Glu Lys His Arg Leu Pro Val Ser Leu Ser Ser Ala Lys Gln
140 145 150

Arg Gln Leu Tyr Lys Asn Arg Gly Phe Leu Pro Leu Ser His Phe 155 160 165

Leu Pro Met Leu Pro Met Val Pro Glu Glu Pro Glu Asp Leu Arg
170 175 . 180

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Tyr Leu Val Val Ala Leu Gly Tyr His Lys Ala Tyr Gly Phe Ser 20 25 30

Ala Pro Lys Asp Gln Gln Val Val Thr Ala Val Glu Tyr Gln Glu
35 40 45

Ala Ile Leu Ala Cys Lys Thr Pro Lys Lys Thr Val Ser Ser Arg
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Leu Glu Trp Lys Lys Leu Gly Arg Ser Val Ser Phe Val Tyr Tyr
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<211> 312

<212> PRT

<213> Homo Sapien

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<223> Synthetic Oligonucleotide Probe

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<210> 66

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<213> Homo Sapien

<400> 69

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Arg Leu Cys Thr Cys Glu Ile Arg Pro Trp Phe Thr Pro Arg Ser 35 40 45

Ile Tyr Met Glu Ala Ser Thr Val Asp Cys Asn Asp Leu Gly Leu
50 55 60

Leu Thr Phe Pro Ala Arg Leu Pro Ala Asn Thr Gln Ile Leu Leu 65 70 75

Leu Gln Thr Asn Asn Ile Ala Lys Ile Glu Tyr Ser Thr Asp Phe
80 85 90

Pro Val Asn Leu Thr Gly Leu Asp Leu Ser Gln Asn Asn Leu Ser 95 100 105

Ser Val Thr Asn Ile Asn Val Lys Lys Met Pro Gln Leu Leu Ser 110 115 120

Val Tyr Leu Glu Glu Asn Lys Leu Thr Glu Leu Pro Glu Lys Cys 125 130 135

Leu Ser Glu Leu Ser Asn Leu Gln Glu Leu Tyr Ile Asn His Asn 140 145 150

Leu Leu Ser Thr Ile Ser Pro Gly Ala Phe Ile Gly Leu His Asn 155 160 165

Leu Leu Arg Leu His Leu Asn Ser Asn Arg Leu Gln Met Ile Asn 170 175 180

Ser Lys Trp Phe Asp Ala Leu Pro Asn Leu Glu Ile Leu Met Ile 185 190 190

Gly Glu Asn Pro Ile Ile Arg Ile Lys Asp Met Asn Phe Lys Pro

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Ile	Ser	Phe	Tyr	Asp 245	Asn	Arg	Leu	Ile	Lys 250	Val	Pro	His	Val	Ala 255
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Lys	Glu	Ile	Ser	Ile 365	His	Ser	Asn	Pro	Ile 370	Arg	Cys	Asp	Cys	Val 375
Ile	Arg	Trp	Met	Asn 380	Met	Asn	Lys	Thr	Asn 385	Ile	Arg	Phe	Met	Glu 390
Pro	Asp	Ser	Leu	Phe 395	Cys	Val	Asp	Pro	Pro 400	Glu	Phe	Gln	Gly	Gln 405
Asn	Val	Arg	Gln	Val 410	His	Phe	Arg	Asp	Met 415	Met	Glu	Ile	Cys	Leu 420
Pro	Leu	Ile	Ala	Pro 425	Glu	Ser	Phe	Pro	Ser 430	Asn	Leu	Asn	Val	Glu 435
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Pro	Asn	Thr	Leu	Thr 470	Asp	Lys	Phe	Tyr	Val 475	His	Ser	Glu	Gly	Thr 480
Leu	Asp	Ile	Asn	Gly 485	Val	Thr	Pro	Lys	Glu 490	Gly	Gly	Leu	Tyr	Thr 495

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Cys Ile Ala Thr Asn Leu Val Gly Ala Asp Leu Lys Ser Val Met Ile Lys Val Asp Gly Ser Phe Pro Gln Asp Asn Asn Gly Ser Leu 515 520 Asn Ile Lys Ile Arg Asp Ile Gln Ala Asn Ser Val Leu Val Ser 530 Trp Lys Ala Ser Ser Lys Ile Leu Lys Ser Ser Val Lys Trp Thr 550 Ala Phe Val Lys Thr Glu Asn Ser His Ala Ala Gln Ser Ala Arg Ile Pro Ser Asp Val Lys Val Tyr Asn Leu Thr His Leu Asn Pro 575 Ser Thr Glu Tyr Lys Ile Cys Ile Asp Ile Pro Thr Ile Tyr Gln Lys Asn Arg Lys Lys Cys Val Asn Val Thr Thr Lys Gly Leu His Pro Asp Gln Lys Glu Tyr Glu Lys Asn Asn Thr Thr Leu Met 625 Ala Cys Leu Gly Gly Leu Leu Gly Ile Ile Gly Val Ile Cys Leu Ile Ser Cys Leu Ser Pro Glu Met Asn Cys Asp Gly Gly His Ser 655 Tyr Val Arg Asn Tyr Leu Gln Lys Pro Thr Phe Ala Leu Gly Glu 665 670 Leu Tyr Pro Pro Leu Ile Asn Leu Trp Glu Ala Gly Lys Glu Lys 680 685 Ser Thr Ser Leu Lys Val Lys Ala Thr Val Ile Gly Leu Pro Thr 700 705 695

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- <212> PRT
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- Ala Ser Met Cys Pro Lys Gly Cys Leu Cys Ser Ser Ser Gly Gly
 35 40 45

Leu Asn Val Thr Cys Ser Asn Ala Asn Leu Lys Glu Ile Pro Arg 50 Asp Leu Pro Pro Glu Thr Val Leu Leu Tyr Leu Asp Ser Asn Gln Ile Thr Ser Ile Pro Asn Glu Ile Phe Lys Asp Leu His Gln Leu 80 85 Arg Val Leu Asn Leu Ser Lys Asn Gly Ile Glu Phe Ile Asp Glu His Ala Phe Lys Gly Val Ala Glu Thr Leu Gln Thr Leu Asp Leu 110 Ser Asp Asn Arg Ile Gln Ser Val His Lys Asn Ala Phe Asn Asn 130 Leu Lys Ala Arg Ala Arg Ile Ala Asn Asn Pro Trp His Cys Asp Cys Thr Leu Gln Gln Val Leu Arg Ser Met Ala Ser Asn His Glu 155 Thr Ala His Asn Val Ile Cys Lys Thr Ser Val Leu Asp Glu His 170 175 Ala Gly Arg Pro Phe Leu Asn Ala Ala Asn Asp Ala Asp Leu Cys 195 Asn Leu Pro Lys Lys Thr Thr Asp Tyr Ala Met Leu Val Thr Met Phe Gly Trp Phe Thr Met Val Ile Ser Tyr Val Val Tyr Tyr Val Arg Gln Asn Gln Glu Asp Ala Arg Arg His Leu Glu Tyr Leu Lys 230 240 Ser Leu Pro Ser Arg Gln Lys Lys Ala Asp Glu Pro Asp Asp Ile

Ser Thr Val Val

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<211> 2290

<212> DNA

<213> Homo Sapien

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35 40 45

Cys Glu Cys Ser Ala Gln Asp Arg Ala Val Leu Cys His Arg Lys

Cys Phe Val Ala Val Pro Glu Gly Ile Pro Thr Glu Thr Arg Leu 65 70 75

Leu Asp Leu Gly Lys Asn Arg Ile Lys Thr Leu Asn Gln Asp Glu 80 85 90

Phe Ala Ser Phe Pro His Leu Glu Glu Leu Glu Leu Asn Glu Asn 95 100 105

Ile Val Ser Ala Val Glu Pro Gly Ala Phe Asn Asn Leu Phe Asn 110 115 120

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<211> 620

<212> PRT

<213> Homo Sapien

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Tyr	Ile	Ser	His	Arg 185	Ala	Phe	Ser	Gly	Leu 190	Asn	Ser	Leu	Glu	Gln 195
Leu	Thr	Leu	Glu	Lys 200	Cys	Asn	Leu	Thr	Ser 205	Ile	Pro	Thr	Glu	Ala 210
Leu	Ser	His	Leu	His 215	Gly	Leu	Ile	Val	Leu 220	Arg	Leu	Arg	His	Leu 225
Asn	Ile	Asn	Ala	Ile 230	Arg	Asp	Tyr	Ser	Phe 235	Lys	Arg	Leu	Tyr	Arg 240
Leu	Lys	Val	Leu	Glu 245	Ile	Ser	His	Trp	Pro 250	Tyr	Leu	Asp	Thr	Met 255
Thr	Pro	Asn	Cys	Leu 260	Tyr	Gly	Leu	Asn	Leu 265	Thr	Ser	Leu	Ser	Ile 270
Thr	His	Cys	Asn	Leu 275	Thr	Ala	Val	Pro	Tyr 280	Leu	Ala	Val	Arg	His 285
Leu	Val	Tyr	Leu	Arg 290	Phe	Leu	Asn	Leu	Ser 295	Tyr	Asn	Pro	Ile	Ser 300
Thr	T1.	a 1	~1											
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			Val	305					310					315
Ile	Gln	Leu	-	305 Gly 320	Gly	Gln	Leu	Ala	310 Val 325	Val	Glu	Pro	Tyr	315 Ala 330
Ile Phe	Gln Arg	Leu Gly	Val	305 Gly 320 Asn 335	Gly Tyr	Gln Leu	Leu Arg	Ala Val	310 Val 325 Leu 340	Val Asn	Glu · Val	Pro Ser	Tyr Gly	315 Ala 330 Asn 345
Ile Phe Gln	Gln Arg Leu	Leu Gly Thr	Val Leu	305 Gly 320 Asn 335 Leu 350	Gly Tyr Glu	Gln Leu Glu	Leu Arg Ser	Ala Val Val	310 Val 325 Leu 340 Phe 355	Val Asn His	Glu Val Ser	Pro Ser Val	Tyr Gly Gly	315 Ala 330 Asn 345 Asn 360
Ile Phe Gln Leu	Gln Arg Leu Glu	Leu Gly Thr	Val Leu Thr	305 Gly 320 Asn 335 Leu 350 Ile 365	Gly Tyr Glu Leu	Gln Leu Glu Asp	Leu Arg Ser	Ala Val Val Asn	310 Val 325 Leu 340 Phe 355 Pro 370	Val Asn His Leu	Glu Val Ser	Pro Ser Val Cys	Tyr Gly Gly Asp	315 Ala 330 Asn 345 Asn 360 Cys 375
Ile Phe Gln Leu Arg	Gln Arg Leu Glu Leu	Leu Gly Thr Thr	Val Leu Thr	305 Gly 320 Asn 335 Leu 350 Ile 365 Val 380	Gly Tyr Glu Leu Phe	Gln Leu Glu Asp	Leu Arg Ser Ser	Ala Val Val Asn	310 Val 325 Leu 340 Phe 355 Pro 370 Trp 385	Val Asn His Leu	Glu Val Ser Ala	Pro Ser Val Cys Asn	Tyr Gly Gly Asp	315 Ala 330 Asn 345 Asn 360 Cys 375 Asn 390
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Asp Pro Pro Pro Ala Ile Leu Trp Leu Ser Pro Arg Lys His Leu 455 460 Val Ser Ala Lys Ser Asn Gly Arg Leu Thr Val Phe Pro Asp Gly Thr Leu Glu Val Arg Tyr Ala Gln Val Gln Asp Asn Gly Thr Tyr Leu Cys Ile Ala Ala Asn Ala Gly Gly Asn Asp Ser Met Pro Ala His Leu His Val Arg Ser Tyr Ser Pro Asp Trp Pro His Gln Pro 515 Asn Lys Thr Phe Ala Phe Ile Ser Asn Gln Pro Gly Glu Gly Glu 530 Ala Asn Ser Thr Arg Ala Thr Val Pro Phe Pro Phe Asp Ile Lys 545 Thr Leu Ile Ile Ala Thr Thr Met Gly Phe Ile Ser Phe Leu Gly Val Val Leu Phe Cys Leu Val Leu Phe Leu Trp Ser Arg Gly 575 Lys Gly Asn Thr Lys His Asn Ile Glu Ile Glu Tyr Val Pro Arg 590 Lys Ser Asp Ala Gly Ile Ser Ser Ala Asp Ala Pro Arg Lys Phe 605 Asn Met Lys Met Ile 620 <210> 74 <211> 22 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 74 tcacctggag cctttattgg cc 22 <210> 75 <211> 23 <212> DNA

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Val Val Leu Lys Cys Gln Val Lys Asp His Glu Asp Ser Ser Leu
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Gln Trp Ser Asn Pro Ala Gln Gln Thr Leu Tyr Phe Gly Glu Lys
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His Glu Leu Ser Ile Ser Ile Ser Asn Val Ala Leu Ala Asp Glu

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<211> 398

<212> PRT

<213> Homo Sapien

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Thr	Gly	Tyr	Lys	Ser 140	Ser	Leu	Arg	Glu	Lys 145	Asp	Thr	Ala	Thr	Leu 150
Asn	Cys	Gln	Ser	Ser 155	Gly	Ser	Lys	Pro	Ala 160	Ala	Arg	Leu	Thr	Trp 165
Arg	Lys	Gly	Asp	Gln 170	Glu	Leu	His	Gly	Glu 175	Pro	Thr	Arg	Ile	Gln 180
Glu	Asp	Pro	Asn	Gly 185	Lys	Thr	Phe	Thr	Val 190	Ser	Ser	Ser	Val	Thr 195
Phe	Gln	Val	Thr	Arg 200	Glu	Asp	Asp	Gly	Ala 205	Ser	Ile	Val	Cys	Ser 210
Val	Asn	His	Glu	Ser 215	Leu	Lys	Gly	Ala	Asp 220	Arg	Ser	Thr	Ser	Gln 225
Arg	Ile	Glu	Val	Leu 230	Tyr	Thr	Pro	Thr	Ala 235	Met	Ile	Arg	Pro	Asp 240
Pro	Pro	His	Pro	Arg 245	Glu	Gly	Gln	Lys	Leu 250	Leu	Leu	His	Cys	Glu 255
Gly	Arg	Gly	Asn	Pro 260	Val	Pro	Gln	Gln	Tyr 265	Leu	Trp	Glu	Lys	Glu 270
Gly	Ser	Val	Pro	Pro 275	Leu	Lys	Met	Thr	Gln 280	Glu	Ser	Ala	Leu	Ile 285
Phe	Pro	Phe	Leu	Asn 290	Lys	Ser	Asp	Ser	Gly 295	Thr	Tyr	Gly	Cys	Thr 300
Ala	Thr	Ser	Asn	Met 305	Gly	Ser	Tyr	Lys	Ala 310	Tyr	Tyr	Thr	Leu	Asn 315
Val	Asn	Asp	Pro	Ser 320	Pro	Val	Pro	Ser	Ser 325	Ser	Ser	Thr	Tyr	His 330
Ala	Ile	Ile	Gly	Gly 335	Ile	Val	Ala	Phe	Ile 340	Val	Phe	Leu	Leu	Leu 345
Ile	Met	Leu	Ile	Phe 350	Leu	Gly	His	Tyr	Leu 355	Ile	Arg	His	Lys	Gly 360
Thr	Tyr	Leu	Thr	His 365	Glu	Ala	Lys	Gly	Ser 370	Asp	Asp	Ala	Pro	Asp 375
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Cys Asn Glu Ile Glu Gly Asp Leu His Val Asp Cys Glu Lys Lys
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Gly Phe Thr Ser Leu Gln Arg Phe Thr Ala Pro Thr Ser Gln Phe
50 55 60

Tyr His Leu Phe Leu His Gly Asn Ser Leu Thr Arg Leu Phe Pro
65 70 75

Asn Glu Phe Ala Asn Phe Tyr Asn Ala Val Ser Leu His Met Glu
80 85 90

Asn Asn Gly Leu His Glu Ile Val Pro Gly Ala Phe Leu Gly Leu 95 100 105

Gln Leu Val Lys Arg Leu His Ile Asn Asn Asn Lys Ile Lys Ser 110 115 120

Phe Arg Lys Gln Thr Phe Leu Gly Leu Asp Asp Leu Glu Tyr Leu 125 130 135

Gln Ala Asp Phe Asn Leu Leu Arg Asp Ile Asp Pro Gly Ala Phe
140 145 150

Gln Asp Leu Asn Lys Leu Glu Val Leu Ile Leu Asn Asp Asn Leu 155 160 165

Ile Ser Thr Leu Pro Ala Asn Val Phe Gln Tyr Val Pro Ile Thr 170 175 180

His Leu Asp Leu Arg Gly Asn Arg Leu Lys Thr Leu Pro Tyr Glu 185 190 195

Glu Val Leu Glu Gln Ile Pro Gly Ile Ala Glu Ile Leu Leu Glu
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Asp Asn Pro Trp Asp Cys Thr Cys Asp Leu Leu Ser Leu Lys Glu 215 220 225

Trp Leu Glu Asn Ile Pro Lys Asn Ala Leu Ile Gly Arg Val Val
230 235 240

Cys Glu Ala Pro Thr Arg Leu Gln Gly Lys Asp Leu Asn Glu Thr
245 250 250

Thr Glu Gln Asp Leu Cys Pro Leu Lys Asn Arg Val Asp Ser Ser 260 265 270

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Thr E	?he	Asn	Ala	Met 470	Pro	Lys	Leu	Arg	Ile 475	Leu	Ile	Leu	Asn	Asn 480
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Leu S									400					
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- Cys Ser Ala Ser Gly Ala Cys Tyr Ser Leu His His Ala Thr Met
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- Lys Arg Gln Ala Ala Glu Glu Ala Cys Ile Leu Arg Gly Gly Ala
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- Leu Ser Thr Val Arg Ala Gly Ala Glu Leu Arg Ala Val Leu Ala 65 70 75

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Gly	Val	Glu	Pro	Ala 155	Gly	Trp	Lys	Glu	Met 160	Arg	Cys	His	Leu	Arg 165
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Ala	Pro	Arg	Pro	Gly 185	Ala	Ala	Ser	Asn	Leu 190	Ser	Tyr	Arg	Ala	Pro 195
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Glu	Val	Ser	Ala	Leu 215	Cys	Arg	Gly	Gln	Leu 220	Pro	Ile	Ser	Val	Thr 225
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Ala	Ile	Phe	Ile	Gly 395	Ala	Val	Ala	Ala	Met 400	Thr	Gly	Tyr	Trp	Leu 405
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- <211> 366
- <212> PRT
- <213> Homo Sapien
- <400> 114
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- Ala Ser Pro Gly Ala Ala Leu Thr Gly Glu Gln Leu Leu Gly Ser
 20 25 30

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Ala	Asp	Met	Glu	Glu 50	Leu	Val	Ile	Pro	Thr 55	His	Val	Arg	Ala	Gln 60
Tyr	Val	Ala	Leu	Leu 65	Gln	Arg	Ser	His	Gly 70	Asp	Arg	Ser	Arg	Gly 75
Lys	Arg	Phe	Ser	Gln 80	Ser	Phe	Arg	Glu	Val 85	Ala	Gly	Arg	Phe	Leu 90
Ala	Leu	Glu	Ala	Ser 95	Thr	His	Leu	Leu	Val 100	Phe	Gly	Met	Glu	Gln 105
Arg	Leu	Pro	Pro	Asn 110	Ser	Glu	Leu	Val	Gln 115	Ala	Val	Leu	Arg	Leu 120
Phe	Gln	Glu	Pro	Val 125	Pro	Lys	Ala	Ala	Leu 130	His	Arg	His	Gly	Arg 135
Leu	Ser	Pro	Arg	Ser 140	Ala	Arg	Ala	Arg	Val 145	Thr	Val	Glu	Trp	Leu 150
Arg	Val	Arg	Asp	Asp 155	Gly	Ser	Asn	Arg	Thr 160	Ser	Leu	Ile	Asp	Ser 165
Arg	Leu	Val	Ser	Val 170	His	Glu	Ser	Gly	Trp 175	Lys	Ala	Phe	Asp	Val 180
Thr	Glu	Ala	Val	Asn 185	Phe	Trp	Gln	Gln	Leu 190	Ser	Arg	Pro	Arg	Gln 195
Pro	Leu	Leu	Leu	Gln 200	Val	Ser	Val	Gln	Arg 205	Glu	His	Leu	Gly	Pro 210
Leu	Ala	Ser	Gly	Ala 215	His	Lys	Leu	Val	Arg 220	Phe	Ala	Ser	Gln	Gly 225
Ala	Pro	Ala	Gly	Leu 230	Gly	Glu	Pro	Gln	Leu 235	Glu	Leu	His	Thr	Leu 240
Asp	Leu	Gly	Asp	Tyr 245	Gly	Àla	Gln	Gly	Asp 250	Cys	Asp	Pro	Glu	Ala 255
Pro	Met	Thr	Glu	Gly 260	Thr	Arg	ĊAa	Cys	Arg 265	Gln	Glu,	Met	Tyr	Ile 270
Asp	Leu	Gln	Gly	Met 275	Lys	Trp	Ala	Glu	Asn 280	Trp	Val	Leu	Glu	Pro 285
Pro	Gly	Phe	Leu	Ala 290	Tyr	Glu	Cys	Val	Gly 295	Thr	Cys	Arg	Gln	Pro 300
Pro	Glu	Ala	Leu	Ala 305	Phe _.	Lys	Trp	Pro	Phe 310	Leu	Gly	Pro	Arg	Gln 315
Cys	Ile	Ala	Ser	Glu 320	Thr	Asp	Ser	Leu	Pro 325	Met	Ile	Val	Ser	Ile 330

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gaggaccggg tgaccttctt gccaactggt atcaccttca agtccgtgac 350 acgggaagac actgggacat acacttgtat ggtctctgag gaaggcggca 400 acagetatgg ggaggteaag gteaagetea tegtgettgt geeteeatee 450 aageetacag ttaacateee eteetetgee accattggga accgggeagt 500 gctgacatgc tcagaacaag atggttcccc accttctgaa tacacctggt 550 tcaaagatgg gatagtgatg cctacgaatc ccaaaagcac ccgtgccttc 600 agcaactctt cctatgtcct gaatcccaca acaggagagc tggtctttga 650 tcccctgtca gcctctgata ctggagaata cagctgtgag gcacggaatg 700 ggtatgggac acccatgact tcaaatgctg tgcgcatgga agctgtggag 750 cggaatgtgg gggtcatcgt ggcagccgtc cttgtaaccc tgattctcct 800 gggaatcttg gtttttggca tctggtttgc ctatagccga ggccactttg 850 acagaacaaa gaaagggact tcgagtaaga aggtgattta cagccagcct 900 agtgcccgaa gtgaaggaga attcaaacag acctcgtcat tcctggtgtg 950 agcctggtcg gctcaccgcc tatcatctgc atttgcctta ctcaggtgct 1000 accggactet ggcccctgat gtctgtagtt tcacaggatg ccttatttgt 1050 cttctacacc ccacagggcc ccctacttct tcggatgtgt ttttaataat 1100 gtcagctatg tgccccatcc tccttcatgc cctccctccc tttcctacca 1150 ctgctgagtg gcctggaact tgtttaaagt gtttattccc catttctttg 1200 agggatcagg aaggaatcct gggtatgcca ttgacttccc ttctaagtag 1250 acagcaaaaa tggcgggggt cgcaggaatc tgcactcaac tgcccacctg 1300 gctggcaggg atctttgaat aggtatcttg agcttggttc tgggctcttt 1350 ccttgtgtac tgacgaccag ggccagctgt tctagagcgg gaattagagg 1400 ctagagegge tgaaatggtt gtttggtgat gacactgggg teetteeate 1450 tetggggeee actetettet gtetteeeat gggaagtgee actgggatee 1500 ctctgccctg tcctcctgaa tacaagctga ctgacattga ctgtgtctgt 1550 ggaaaatggg agctcttgtt gtggagagca tagtaaattt tcagagaact 1600 tgaagccaaa aggatttaaa accgctgctc taaagaaaag aaaactggag 1650 gctgggcgca gtggctcacg cctgtaatcc cagaggctga ggcaggcgga 1700 tcacctgagg tcgggagttc gggatcagcc tgaccaacat ggagaaaccc 1750

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- <210> 119
- <211> 299
- <212> PRT
- <213> Homo Sapien

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- Ile Leu Ala Ile Leu Cys Ser Leu Ala Leu Gly Ser Val Thr
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- Val His Ser Ser Glu Pro Glu Val Arg Ile Pro Glu Asn Asn Pro 35 40 45
- Val Lys Leu Ser Cys Ala Tyr Ser Gly Phe Ser Ser Pro Arg Val
 50 55 60
- Glu Trp Lys Phe Asp Gln Gly Asp Thr Thr Arg Leu Val Cys Tyr
 65 70 75
- Asn Asn Lys Ile Thr Ala Ser Tyr Glu Asp Arg Val Thr Phe Leu 80 85 90
- Pro Thr Gly Ile Thr Phe Lys Ser Val Thr Arg Glu Asp Thr Gly
 95 100 105
- Thr Tyr Thr Cys Met Val Ser Glu Glu Gly Gly Asn Ser Tyr Gly 110 115 120
- Glu Val Lys Val Lys Leu Ile Val Leu Val Pro Pro Ser Lys Pro 125 130 135
- Thr Val Asn Ile Pro Ser Ser Ala Thr Ile Gly Asn Arg Ala Val 140 145 150
- Leu Thr Cys Ser Glu Gln Asp Gly Ser Pro Pro Ser Glu Tyr Thr 155 160 165
- Trp Phe Lys Asp Gly Ile Val Met Pro Thr Asn Pro Lys Ser Thr 170 175 180
- Arg Ala Phe Ser Asn Ser Ser Tyr Val Leu Asn Pro Thr Thr Gly
 185 190 195
- Glu Leu Val Phe Asp Pro Leu Ser Ala Ser Asp Thr Gly Glu Tyr 200 205 210
- Ser Cys Glu Ala Arg Asn Gly Tyr Gly Thr Pro Met Thr Ser Asn 215 220 225
- Ala Val Arg Met Glu Ala Val Glu Arg Asn Val Gly Val Ile Val 230 235 240

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Ala Ala Val Leu Val Thr Leu Ile Leu Leu Gly Ile Leu Val Phe
 Gly Ile Trp Phe Ala Tyr Ser Arg Gly His Phe Asp Arg Thr Lys
 Lys Gly Thr Ser Ser Lys Lys Val Ile Tyr Ser Gln Pro Ser Ala
Arg Ser Glu Gly Glu Phe Lys Gln Thr Ser Ser Phe Leu Val
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<213> Homo Sapien
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gcgcaggttg gagcgtggcg aacaggggct ctgggcctgg cgctgctgct 100
gctgctcggc ctcggactag gcctggaggc cgccgcgagc ccgctttcca 150
ccccgacctc tgcccaggcc gcaggcccca gctcaggctc gtgcccaccc 200
accaagttcc agtgccgcac cagtggctta tgcgtgcccc tcacctggcg 250
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gagcgcctcc gcccactggg gttactggtg gccatgaagg agtccctgct 850
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- <211> 282
- <212> PRT
- <213> Homo Sapien

<400> 127

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- Leu Glu Ala Ala Ser Pro Leu Ser Thr Pro Thr Ser Ala Gln
 35 40 45
- Ala Ala Gly Pro Ser Ser Gly Ser Cys Pro Pro Thr Lys Phe Gln 50 55 60
- Cys Arg Thr Ser Gly Leu Cys Val Pro Leu Thr Trp Arg Cys Asp
 65 70 75
- Arg Asp Leu Asp Cys Ser Asp Gly Ser Asp Glu Glu Glu Cys Arg 80 85 90
- Ile Glu Pro Cys Thr Gln Lys Gly Gln Cys Pro Pro Pro Gly 95 100 105
- Leu Pro Cys Pro Cys Thr Gly Val Ser Asp Cys Ser Gly Gly Thr
 110 115 120
- Asp Lys Lys Leu Arg Asn Cys Ser Arg Leu Ala Cys Leu Ala Gly 125 130 135
- Glu Leu Arg Cys Thr Leu Ser Asp Asp Cys Ile Pro Leu Thr Trp
 140 145 150
- Arg Cys Asp Gly His Pro Asp Cys Pro Asp Ser Ser Asp Glu Leu 155 160 165
- Gly Cys Gly Thr Asn Glu Ile Leu Pro Glu Gly Asp Ala Thr Thr 170 175 180
- Met Gly Pro Pro Val Thr Leu Glu Ser Val Thr Ser Leu Arg Asn

				185					190					195
Ala '	Thr	Thr	Met	Gly 200	Pro	Pro	Val	Thr	Leu 205	Glu	Ser	Val	Pro	Ser 210
Val (Gly	Asn	Ala	Thr 215	Ser	Ser	Ser	Ala	Gly 220	Asp	Gln	Ser	Gly	Ser 225
Pro '	Thr	Ala	Tyr	Gly 230	Val	Ile	Ala	Ala	Ala 235	Ala	Val	Leu	Ser	Ala 240
Ser :	Leu	Val	Thr	Ala 245	Thr	Leu	Leu	Leu	Leu 250	Ser	Trp	Leu	Arg	Ala 255
Gln (Glu	Arg	Leu	Arg 260	Pro	Leu	Gly	Leu	Leu 265	Val	Ala	Met	Lys	Glu 270
Ser :	Leu	Leu	Leu	Ser 275	Glu	Gln	Lys	Thr	Ser 280	Leu	Pro			
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<210> 132

<211> 490

<212> PRT

<213> Homo Sapien

<400> 132

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Gln Leu Thr Gly Gly Phe Asp Asp Leu Gln Val Cys Ala Asp Pro 50 55 60

Gly Ile Pro Glu Asn Gly Phe Arg Thr Pro Ser Gly Gly Val Phe
65 70 75

Phe Glu Gly Ser Val Ala Arg Phe His Cys Gln Asp Gly Phe Lys 80 85 90

Leu Lys Gly Ala Thr Lys Arg Leu Cys Leu Lys His Phe Asn Gly
95 100 105

Thr Leu Gly Trp Ile Pro Ser Asp Asn Ser Ile Cys Val Glu 110 115 120

Asp Cys Arg Ile Pro Gln Ile Glu Asp Ala Glu Ile His Asn Lys 125 130 135

Thr Tyr Arg His Gly Glu Lys Leu Ile Ile Thr Cys His Glu Gly
140 145 150

Phe Lys Ile Arg Tyr Pro Asp Leu His Asn Met Val Ser Leu Cys 155 160 165

Arg Asp Asp Gly Thr Trp Asn Asn Leu Pro Ile Cys Gln Gly Cys
170 175 180

Leu	Arg	Pro	Leu	Ala 185	Ser	Ser	Asn	Gly	Tyr 190	Val	Asn	Ile	Ser	Glu 195
Leu	Gln	Thr	Ser	Phe 200	Pro	Val	Gly	Thr	Val 205	Ile	Ser	Tyr	Arg	Cys 210
Phe	Pro	Gly	Phe	Lys 215	Leu	Asp	Gly	Ser	Ala 220	Tyr	Leu	Glu	Cys	Leu 225
Gln	Asn	Leu	Ile	Trp 230	Ser	Ser	Ser	Pro	Pro 235	Arg	Cys	Leu	Ala	Leu 240
Glu	Ala	Gln	Val	Cys 245	Pro	Leu	Pro	Pro	Met 250	Val	Ser	His	Gly	Asp 255
Phe	Val	Cys	His	Pro 260	Arg	Pro	Cys	Glu	Arg 265	Tyr	Asn	His	Gly	Thr 270
Val	Val	Glu	Phe	Tyr 275	Cys	Asp	Pro	Gly	Tyr 280	Ser	Leu	Thr	Ser	Asp 285
Tyr	Lys	Tyr	Ile	Thr 290	Cys	Gln	Tyr	Gly	Glu 295	Trp	Phe	Pro	Ser	Tyr 300
Gln	Val	Tyr	Cys	Ile 305	Lys	Ser	Glu	Gln	Thr 310	Trp	Pro	Ser	Thr	His 315
Glu	Thr	Leu	Leu	Thr 320	Thr	Trp	Lys	Ile	Val 325	Ala	Phe	Thr	Ala	Thr 330
Ser	Val	Leu	Leu	Val 335	Leu	Leu	Leu	Val	Ile 340	Leu	Ala	Arg	Met	Phe 345
Gln	Thr	Lys	Phe	Lys 350	Ala	His	Phe	Pro	Pro 355	Arg	Gly	Pro	Pro	Arg 360
Ser	Ser	Ser	Ser	Asp 365	Pro	Asp	Phe	Val	Val 370	Val	Asp	Gly	Val	Pro 375
Val	Met	Leu	Pro	Ser 380	Tyr	Asp	Glu	Ala	Val 385	Ser	Gly	Gly	Leu	Ser 390
Ala	Leu	Gly	Pro	Gly 395	Tyr	Met	Ala	Ser	Val 400	Gly	Gln	Gly	Cys	Pro 405
Leu	Pro	Val	Asp	Asp 410	Gln	Ser	Pro	Pro	Ala 415	Tyr	Pro	Gly	Ser	Gly 420
Asp	Thr	Asp	Thr	Gly 425	Pro	Gly	Glu	Ser	Glu 430	Thr	Cys	Asp	Ser	Val 435
Ser	Gly	Ser	Ser	Glu 440	Leu	Leu	Gln	Ser	Leu 445	Tyr	Ser	Pro	Pro	Arg 450
Cys	Gln	Glu	Ser	Thr 455	His	Pro	Ala	Ser	Asp 460	Asn	Pro	Asp	Ile	Ile 465
Ala	Ser	Thr	Ala	Glu	Glu	Val	Ala	Ser	Thr	Ser	Pro	Gly	Ile	His

His Ala His Trp Val Leu Phe Leu Arg Asn 485 490

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- <211> 23
- <212> DNA
- <213> Artificial Sequence
- <220>
- <223> Synthetic Oligonucleotide Probe
- <400> 133

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- <210> 134
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- <212> DNA
- <213> Artificial Sequence
- <220>
- <223> Synthetic Oligonucleotide Probe
- <400> 134

agccaggatc gcagtaaaac tcc 23

- <210> 135
- <211> 50
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- <213> Artificial Sequence
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- <400> 135

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- <210> 136
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- <212> DNA
- <213> Homo Sapien
- <400> 136
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- cccgccagcc cgctccaccg ccgtagcgcc cgagtgtcgg ggggcgcacc 150
- cgagtcgggc catgaggccg ggaaccgcgc tacaggccgt gctgctggcc 200
- gtgctgctgg tggggctgcg ggccgcgacg ggtcgcctgc tgagtgcctc 250
- ggatttggac ctcagaggag ggcagccagt ctgccgggga gggacacaga 300
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<212> PRT

<213> Homo Sapien

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Thr Lys Lys Gln His Thr Ile Trp Pro Ser Pro His Gln Gly Asn Ser Pro Asp Leu Glu Val Tyr Asn Val Ile Arg Lys Gln Ser Glu 290 295 Ala Asp Leu Ala Glu Thr Arg Pro Asp Leu Lys Asn Ile Ser Phe Arg Val Cys Ser Gly Glu Ala Thr Pro Asp Asp Met Ser Cys Asp Tyr Asp Asn Met Ala Val Asn Pro Ser Glu Ser Gly Phe Val Thr Leu Val Ser Val Glu Ser Gly Phe Val Thr Asn Asp Ile Tyr Glu Phe Ser Pro Asp Gln Met Gly Arg Ser Lys Glu Ser Gly Trp Val 370 Glu Asn Glu Ile Tyr Gly Tyr <210> 138 <211> 50 <212> DNA <213> Artificial Sequence <223> Synthetic Oligonucleotide Probe <400> 138 gttcattgaa aacctcttgc catctgatgg tgacttctgg attgggctca 50 <210> 139 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 139 aagccaaaga agcctgcagg aggg 24 <210> 140 <211> 24 <212> DNA <213> Artificial Sequence <223> Synthetic Oligonucleotide Probe <400> 140 cagtccaagc ataaaggtcc tggc 24 <210> 141 <211> 1514

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- <210> 142
- <211> 428
- <212> PRT
- <213> Homo Sapien

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- Val Leu Thr Ser Leu Ala Tyr Cys Leu His Gln Arg Arg Val Ala 20 25 30
- Leu Ala Glu Leu Gln Glu Ala Asp Gly Gln Cys Pro Val Asp Arg
 35 40 45
- Ser Leu Leu Lys Leu Lys Met Val Gln Val Val Phe Arg His Gly 50 55 60
- Ala Arg Ser Pro Leu Lys Pro Leu Pro Leu Glu Glu Gln Val Glu 65 70 75
- Trp Asn Pro Gln Leu Leu Glu Val Pro Pro Gln Thr Gln Phe Asp 80 85 90
- Tyr Thr Val Thr Asn Leu Ala Gly Gly Pro Lys Pro Tyr Ser Pro 95 100 105
- Tyr Asp Ser Gln Tyr His Glu Thr Thr Leu Lys Gly Gly Met Phe 110 115 120
- Ala Gly Gln Leu Thr Lys Val Gly Met Gln Gln Met Phe Ala Leu 125 130 135
- Gly Glu Arg Leu Arg Lys Asn Tyr Val Glu Asp Ile Pro Phe Leu 140 145 150
- Ser Pro Thr Phe Asn Pro Gln Glu Val Phe Ile Arg Ser Thr Asn 155 160 165
- Ile Phe Arg Asn Leu Glu Ser Thr Arg Cys Leu Leu Ala Gly Leu
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- Phe Gln Cys Gln Lys Glu Gly Pro Ile Ile Ile His Thr Asp Glu 185 190 195
- Ala Asp Ser Glu Val Leu Tyr Pro Asn Tyr Gln Ser Cys Trp Ser
 200 205 210
- Leu Arg Gln Arg Thr Arg Gly Arg Arg Gln Thr Ala Ser Leu Gln 215 220 225
- Pro Gly Ile Ser Glu Asp Leu Lys Lys Val Lys Asp Arg Met Gly

230 235 240

Ile Asp Ser Ser Asp Lys Val Asp Phe Phe Ile Leu Leu Asp Asn 245 250 255

Val Ala Ala Glu Gln Ala His Asn Leu Pro Ser Cys Pro Met Leu 260 265 270

Lys Arg Phe Ala Arg Met Ile Glu Gln Arg Ala Val Asp Thr Ser 275 280 280

Leu Tyr Ile Leu Pro Lys Glu Asp Arg Glu Ser Leu Gln Met Ala 290 295 300

Val Gly Pro Phe Leu His Ile Leu Glu Ser Asn Leu Leu Lys Ala 305 310 315

Met Asp Ser Ala Thr Ala Pro Asp Lys Ile Arg Lys Leu Tyr Leu
320 325 330

Tyr Ala Ala His Asp Val Thr Phe Ile Pro Leu Leu Met Thr Leu
335 340 345

Gly Ile Phe Asp His Lys Trp Pro Pro Phe Ala Val Asp Leu Thr 350 355 360

Met Glu Leu Tyr Gln His Leu Glu Ser Lys Glu Trp Phe Val Gln 365 370 375

Leu Tyr Tyr His Gly Lys Glu Gln Val Pro Arg Gly Cys Pro Asp 380 385 390

Gly Leu Cys Pro Leu Asp Met Phe Leu Asn Ala Met Ser Val Tyr 395 400 405

Thr Leu Ser Pro Glu Lys Tyr His Ala Leu Cys Ser Gln Thr Gln $410 \,$ $415 \,$ 420

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<211> 45
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<211> 1686
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<213> Homo Sapien
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- <211> 347
- <212> PRT
- <213> Homo Sapien
- <400> 148
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- Gly Phe Leu Ala Ser Pro Ser Gly Val Arg Leu Val Gly Gly Leu
 20 25 30
- His Arg Cys Glu Gly Arg Val Glu Val Glu Gln Lys Gly Gln Trp 35 40 45
- Gly Thr Val Cys Asp Asp Gly Trp Asp Ile Lys Asp Val Ala Val
 50 55 60
- Leu Cys Arg Glu Leu Gly Cys Gly Ala Ala Ser Gly Thr Pro Ser

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Gly	Ile	Leu	Tyr	Glu 80	Pro	Pro	Ala	Glu	Lys 85	Glu	Gln	Lys	Val	Leu 90
Ile	Gln	Ser	Val	Ser 95	Cys	Thr	Gly	Thr	Glu 100	Asp	Thr	Leu	Ala	Gln 105
Cys	Glu	Gln	Glu	Glu 110	Val	Tyr	Asp	Cys	Ser 115	His	Asp	Glu	Asp	Ala 120
Gly	Ala	Ser	Cys	Glu 125	Asn	Pro	Glu	Ser	Ser 130	Phe	Ser	Pŕo	Val	Pro 135
Glu	Gly	Val	Arg	Leu 140	Ala	Asp	Gly	Pro	Gly 145	His	Cys	Lys	Gly	Arg 150
Val	Glu	Val	Lys	His 155	Gln	Asn	Gln	Trp	Tyr 160	Thr	Val	Cys	Gln	Thr 165
Gly	Trp	Ser	Leu	Arg 170	Ala	Ala	Lys	Val	Val 175	Cys	Arg	Gln	Leu	Gly 180
Cys	Gly	Arg	Ala	Val 185	Leu	Thr	Gln	Lys	Arg 190	Cys	Asn	Lys	His	Ala 195
Tyr	Gly	Arg	Lys	Pro 200	Ile	Trp	Leu	Ser	Gln 205	Met	Ser	Cys	Ser	Gly 210
Arg	Glu	Ala	Thr	Leu 215	Gln	Asp	Cys	Pro	Ser 220	Gly	Pro	Trp	Gly	Lys 225
Asn	Thr	Cys	Asn	His 230	Asp	Glu	Asp	Thr	Trp 235	Val	Glu	Cys	Glu	Asp 240
Pro	Phe	Asp	Leu	Arg 245	Leu	Val	Gly	Gly	Asp 250	Asn	Leu	Cys	Ser	Gly 255
Arg	Leu	Glu	Val	Leu 260	His	Lys	Gly	Val	Trp 265	Gly	Ser	Val	Cys	Asp 270
Asp	Asn	Trp	Gly	Glu 275	_	Glu	Asp	Gln	Val 280	Val	Cys	Lys	Gln	Leu 285
Gly	Cys	Gly	Lys	Ser 290	Leu	Ser	Pro	Ser	Phe 295	Arg	Asp	Arg	Lys	Cys 300
Tyr	Gly	Pro	Gly	Val 305	Gly	Arg	Ile	Trp	Leu 310	Asp	Asn	Val	Arg	Cys 315
Ser	Gly	Glu	Glu	Gln 320	Ser	Leu	Glu	Gln	Cys 325	Gln	His	Arg	Phe	Trp 330
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<210> 151
<211> 50
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<211> 1427
<212> DNA
<213> Homo Sapien
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cacagocato otgoccotgo tgttoggotg cotgggogto ttoggootot 200
tccggctgct gcagtgggtg cgcgggaagg cctacctgcg gaatgctgtg 250
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<211> 310

<212> PRT

<213> Homo Sapien

<400> 153

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Gly Lys Ala Tyr Leu Arg Asn Ala Val Val Val Ile Thr Gly Ala 35 40 45

Thr Ser Gly Leu Gly Lys Glu Cys Ala Lys Val Phe Tyr Ala Ala 50 55 60

Gly Ala Lys Leu Val Leu Cys Gly Arg Asn Gly Gly Ala Leu Glu 65 70 75

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Glu Leu Ile Arq Glu Leu Thr Ala Ser His Ala Thr Lys Val Gln
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Thr His Lys Pro Tyr Leu Val Thr Phe Asp Leu Thr Asp Ser Gly
                  95
                                     100
                                                         105
Ala Ile Val Ala Ala Ala Glu Ile Leu Gln Cys Phe Gly Tyr
Val Asp Ile Leu Val Asn Asn Ala Gly Ile Ser Tyr Arg Gly Thr
Ile Met Asp Thr Thr Val Asp Val Asp Lys Arg Val Met Glu Thr
Asn Tyr Phe Gly Pro Val Ala Leu Thr Lys Ala Leu Leu Pro Ser
                 155
Met Ile Lys Arg Arg Gln Gly His Ile Val Ala Ile Ser Ser Ile
Gln Gly Lys Met Ser Ile Pro Phe Arg Ser Ala Tyr Ala Ala Ser
Lys His Ala Thr Gln Ala Phe Phe Asp Cys Leu Arg Ala Glu Met
Glu Gln Tyr Glu Ile Glu Val Thr Val Ile Ser Pro Gly Tyr Ile
His Thr Asn Leu Ser Val Asn Ala Ile Thr Ala Asp Gly Ser Arg
                230
                                     235
Tyr Gly Val Met Asp Thr Thr Ala Gln Gly Arg Ser Pro Val
                245
Glu Val Ala Gln Asp Val Leu Ala Ala Val Gly Lys Lys Lys
                260
                                     265
                                                         270
Asp Val Ile Leu Ala Asp Leu Leu Pro Ser Leu Ala Val Tyr Leu
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Arg Thr Leu Ala Pro Gly Leu Phe Phe Ser Leu Met Ala Ser Arg
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<210> 154
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- <213> Artificial Sequence
- <223> Synthetic Oligonucleotide Probe
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<211> 1771
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ctccctagag tccttcgtga agctttttat tcctaagagg agaaaatcag 200
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ctgactgcct atgaatttgc taaacttaaa agcaagctgg ttctctggga 300
tataaataag catggactgg aggaaacagc tgccaaatgc aagggactgg 350
gtgccaaggt tcataccttt gtggtagact gcagcaaccg agaagatatt 400
tacagctctg caaagaaggt gaaggcagaa attggagatg ttagtatttt 450
agtaaataat gctggtgtag tctatacatc agatttgttt gctacacaag 500
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- <211> 300
- <212> PRT
- <213> Homo Sapien

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His	Gly	Ile	Gly	Arg 50	Leu	Thr	Ala	Tyr	Glu 55	Phe	Ala	ГÀЗ	Leu.	Lys 60
Ser	Lys	Leu	Val	Leu 65	Trp	Asp	Ile	Asn	Lys 70	His	Gly	Leu	Glu	Glu 75
Thr	Ala	Ala	Lys	S0	Lys	Gly	Leu	Gly	Ala 85	Lys	Val	His	Thr	Phe 90
Val	Val	Asp	Cys	Ser 95	Asn	Arg	Glu	Asp	Ile 100	Tyr	Ser	Ser	Ala	Lys 105
Lys	Val	Lys	Ala	Glu 110	Ile	Gly	Asp	Val	Ser 115	Ile	Leu	Val	Asn	Asn 120
Ala	Gly	Val	Val	Tyr 125	Thr	Ser	Asp	Leu	Phe 130	Ala	Thr	Gln	Asp	Pro 135
Gln	Ile	Glu	Lys	Thr 140	Phe	Glu	Val	Asn	Val 145	Leu	Ala	His	Phe	Trp 150
Thr	Thr	Lys	Ala	Phe 155	Leu	Pro	Ala	Met	Thr 160	Lys	Asn	Asn	His	Gly 165
His	Ile	Val	Thr	Val 170	Ala	Ser	Ala	Ala	Gly 175	His	Val	Ser	Val	Pro 180
Phe	Leu	Leu	Ala	Tyr 185	Cys	Ser	Ser	Lys	Phe 190	Ala	Ala	Val	Gly	Phe 195
His	Lys	Thr	Leu	Thr 200	Asp	Glu	Leu	Ala	Ala 205	Leu	Gln	Ile	Thr	Gly 210
Val	Lys	Thr	Thr	Cys 215	Leu	Суѕ	Pro	Asn	Phe 220	Val	Asn	Thr	Gly	Phe 225
Ile	Lys	Asn	Pro	Ser 230	Thr	Ser	Leu	Gly	Pro 235	Thr	Leu	Glu	Pro	Glu 240
Glu	Val	Val	Asn	Arg 245	Leu	Met	His	Gly	Ile 250	Leu	Thr	Glu	Gln	Lys 255
Met	Ile	Phe	Ile	Pro 260	Ser	Ser	Ile	Ala	Phe 265	Leu	Thr	Thr	Leu	Glu 270
Arg	Ile	Leu	Pro	Glu 275	Arg	Phe	Leu	Ala	Val 280	Leu	Lys	Arg	Lys	Ile 285
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Thr Pro Tyr Ile Glu Ala Gly Lys Ile Gln Lys Gly Arg Glu Leu
Ser Leu Val Gly Pro Phe Pro Gly Leu Asn Met Lys Ser Tyr Ala
Gly Phe Leu Thr Val Asn Lys Thr Tyr Asn Ser Asn Leu Phe Phe
Trp Phe Phe Pro Ala Gln Ile Gln Pro Glu Asp Ala Pro Val Val
Leu Trp Leu Gln Gly Gly Pro Gly Gly Ser Ser Met Phe Gly Leu
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Phe Val Glu His Gly Pro Tyr Val Val Thr Ser Asn Met Thr Leu
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Arg Asp Arg Asp Phe Pro Trp Thr Thr Leu Ser Met Leu Tyr
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                                     145
Ile Asp Asn Pro Val Gly Thr Gly Phe Ser Phe Thr Asp Asp Thr
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His Gly Tyr Ala Val Asn Glu Asp Asp Val Ala Arg Asp Leu Tyr
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Ser Ala Leu Ile Gln Phe Phe Gln Ile Phe Pro Glu Tyr Lys Asn
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Asn Asp Phe Tyr Val Thr Gly Glu Ser Tyr Ala Gly Lys Tyr Val
Pro Ala Ile Ala His Leu Ile His Ser Leu Asn Pro Val Arq Glu
Val Lys Ile Asn Leu Asn Gly Ile Ala Ile Gly Asp Gly Tyr Ser
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Asp Pro Ser Tyr Phe Gln Asn Val Thr Gly Cys Ser Asn Tyr Tyr
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 Lys Phe Leu Ser Leu Pro Glu Val Arg Gln Ala Ile His Val Gly
Asn Gln Thr Phe Asn Asp Gly Thr Ile Val Glu Lys Tyr Leu Arg
 Glu Asp Thr Val Gln Ser Val Lys Pro Trp Leu Thr Glu Ile Met
Asn Asn Tyr Lys Val Leu Ile Tyr Asn Gly Gln Leu Asp Ile Ile
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Gly Asp Phe His Gln Val Ile Ile Arg Gly Gly His Ile Leu
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Pro Tyr Asp Gln Pro Leu Arg Ala Phe Asp Met Ile Asn Arg Phe
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- <223> Synthetic Oligonucleotide Probe
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- <210> 166
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<213> Homo Sapien

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Arg Leu Leu Thr Ala Ala Pro Leu Ser Met Glu Gln Arg Gln Pro 50 55 60

Trp Pro Arg Ala Leu Glu Val Asp Ser Arg Ser Val Val Leu Leu
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Ser Val Val Trp Val Leu Leu Ala Pro Pro Ala Ala Gly Met Pro 80 85 90

Gln Phe Ser Thr Phe His Ser Glu Asn Arg Asp Trp Thr Phe Asn 95 100 105

His Leu Thr Val His Gln Gly Thr Gly Ala Val Tyr Val Gly Ala 110 115 120

Ile Asn Arg Val Tyr Lys Leu Thr Gly Asn Leu Thr Ile Gln Val 125 130 135

Ala His Lys Thr Gly Pro Glu Glu Asp Asn Lys Ser Arg Tyr Pro 140 145 150

Pro Leu Ile Val Gln.Pro Cys Ser Glu Val Leu Thr Leu Thr Asn 15.5 160 165

Asn Val Asn Lys Leu Leu Ile Ile Asp Tyr Ser Glu Asn Arg Leu 170 175 180

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Ile	Val	Arg	Ser	Glu 230	Gly	Glu	Asp	Gly	Lys 235	Leu	Phe	Ile	Gly	Thr 240
Ala	Val	Asp	Gly	Lys 245	Gln	Asp	Tyr	Phe	Pro 250	Thr	Leu	Ser	Ser	Arg 255
Lys	Leu	Pro	Arg	Asp 260	Pro	Glu	Ser	Ser	Ala 265	Met	Leu	Asp	Tyr	Glu 270
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Pro	Leu	Gly	Gly	Ser	Thr	Pro	Val	Glu	Gly	Leu	Thr	Leu	Tyr	Thr

470 475 480 Thr Ser Arg Asp Arg Met Thr Ser Val Ala Ser Tyr Val Tyr Asn Gly Tyr Ser Val Val Phe Val Gly Thr Lys Ser Gly Lys Leu Lys 505 Lys Val Arg Val Tyr Glu Phe Arg Cys Ser Asn Ala Ile His Leu Leu Ser Lys Glu Ser Leu Leu Glu Gly Ser Tyr Trp Trp Arg Phe 530 535 Asn Tyr Arg Gln Leu Tyr Phe Leu Gly Glu Gln Arg 545 <210> 171 <211> 20 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 171 tggaataccg cctcctgcag 20 <210> 172 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 172 cttctgccct ttggagaaga tggc 24 <210> 173 <211> 43 <212> DNA <213> Artificial Sequence <223> Synthetic oligonucleotide probe <400> 173 ggactcactg gcccaggcct tcaatatcac cagccaggac gat 43 <210> 174 <211> 3106 <212> DNA <213> Homo Sapien

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<223> unknown base

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attettggae tacaagacaa egaagattge tgteeceetg atceagggtt 1500 acaccgtgct gaggatcttg gtggagaatc gtgggcgagt caactatggg 1550 gagaatattg atgaccagcg caaaggctta attggaaatc tctatctgaa 1600 tgattcaccc ctgaaaaact tcagaatcta tagcctggat atgaagaaga 1650 gettetttea gaggttegge etggacaaat ggngtteeet eecagaaaca 1700 cccacattac ctgctttctt cttgggtagc ttgtccatca gctccacgcc 1750 ttgtgacacc tttctgaagc tggagggctg ggagaagggg gttgtattca 1800 tcaatggcca gaaccttgga cgttactgga acattggacc ccagaagacg 1850 ctttacctcc caggtccctg gttgagcagc ggaatcaacc aggtcatcgt 1900 ttttgaggag acgatggcgg gccctgcatt acagttcacg gaaacccccc 1950 acctgggcag gaaccagtac attaagtgag cggtggcacc ccctcctgct 2000 ggtgccagtg ggagactgcc gcctcctctt gacctgaagc ctggtggctg 2050 ctgccccacc cctcactgca aaagcatctc cttaagtagc aacctcaggg 2100 actgggggct acagtctgcc cctgtctcag ctcaaaaccc taagcctgca 2150 gggaaaggtg ggatggctct gggcctggct ttgttgatga tggctttcct 2200 acagecetge tettgtgeeg aggetgtegg getgteteta gggtgggage 2250 agctaatcag atcgcccagc ctttggccct cagaaaaagt gctgaaacgt 2300 gcccttgcac cggacgtcac agccctgcga gcatctgctg gactcaggcg 2350 tgctctttgc tggttcctgg gaggcttggc cacatccctc atggccccat 2400 tttatccccg aaatcctggg tgtgtcacca gtgtagaggg tggggaaggg 2450 gtgtctcacc tgagctgact ttgttcttcc ttcacaacct tctgagcctt 2500 ctttgggatt ctggaaggaa ctcggcgtga gaaacatgtg acttcccctt 2550 tecetteeca etegetgett eecacagggt gacaggetgg getggagaaa 2600 cagaaatcct caccetgegt cttcccaagt tagcaggtgt ctctggtgtt 2650 cagtgaggag gacatgtgag tcctggcaga agccatggcc catgtctgca 2700 catccaggga ggaggacaga aggcccagct cacatgtgag tcctggcaga 2750 agccatggcc catgtctgca catccaggga ggaggacaga aggcccagct 2800 cacatgtgag tcctggcaga agccatggcc catgtctgca catccaggga 2850 ggaggacaga aggcccagct cacatgtgag tcctggcaga agccatggcc 2900 catgtctgca catccaggga ggaggacaga aggcccagct cagtggcccc 2950

cgctcccac ccccacgcc cgaacagcag gggcagagca gccctccttc 3000 gaagtgtgtc caagtccgca tttgagcctt gttctggggc ccagcccaac 3050 acctggcttg ggctcactgt.cctgagttgc agtaaagcta taaccttgaa 3100 tcacaa 3106

<210> 175

<211> 636

<212> PRT

<213> Homo Sapien

<220>

<221> unsure

<222> 539

<223> unknown amino acid

<400> 175

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Asp Trp Ser Thr Leu Val Pro Leu Arg Leu Arg His Arg Gln Leu
35 40 45

Gly Leu Gln Ala Lys Gly Trp Asn Phe Met Leu Glu Asp Ser Thr
50 55 60

Phe Trp Ile Phe Gly Gly Ser Ile His Tyr Phe Arg Val Pro Arg
65 70 75

Glu Tyr Trp Arg Asp Arg Leu Leu Lys Met Lys Ala Cys Gly Leu 80 85 90

Asn Thr Leu Thr Thr Tyr Val Pro Trp Asn Leu His Glu Pro Glu
95 100 105

Arg Gly Lys Phe Asp Phe Ser Gly Asn Leu Asp Leu Glu Ala Phe 110 115 120

Val Leu Met Ala Ala Glu Ile Gly Leu Trp Val Ile Leu Arg Pro 125 130 135

Gly Pro Tyr Ile Cys Ser Glu Met Asp Leu Gly Gly Leu Pro Ser 140 145 150

Gly Phe Thr Glu Ala Val Asp Leu Tyr Phe Asp His Leu Met Ser

Arg Val Val Pro Leu Gln Tyr Lys Arg Gly Gly Pro Ile Ile Ala 185 190 195

Val Gln Val Glu Asn Glu Tyr Gly Ser Tyr Asn Lys Asp Pro Ala 200 205 210

Tyr	Met	Pro	Tyr	Val 215	Lys	Lys	Ala	Leu	Glu 220	Asp	Arg	Gly	Ile	Val 225
Glu	Leu	Leu	Leu	Thr 230	Ser	Asp	Asn	Lys	Asp 235	Gly	Leu	Ser	Lys	Gly 240
Ile	Val	Gln	Gly	Val 245	Leu	Ala	Thr	Ile	Asn 250	Leu	Gln	Ser	Thr	His 255
Glu	Leu	Gln	Leu	Leu 260	Thr	Thr	Phe	Leu	Phe 265	Asn	Val	Gln	Gly	Thr 270
Gln	Pro	Lys	Met	Val 275	Met	Glu	Tyr	Trp	Thr 280	Gly	Trp	Phe	Asp	Ser 285
Trp	Gly	Gly	Pro	His 290	Asn	Ile	Leu	Asp	Ser 295	Ser	Glu	Val	Leu	Lys 300
Thr	Val	Ser	Ala	Ile 305	Val	Asp	Ala	Gly	Ser 310	Ser	Ile	Asn	Leu	Tyr 315
Met	Phe	His	Gly	Gly 320	Thr	Asn	Phe	Gly	Phe 325	Met	Asn	Gly	Ala	Met 330
His	Phe	His	Asp	Tyr 335	Lys	Ser	Asp	Val	Thr 340	Ser	Tyr	Asp	Tyr	Asp 345
Ala	Val	Leu	Thr	Glu 350	Ala	Gly	Asp	Tyr	Thr 355	Ala	Lys	Tyr	Met	Lys 360
Leu	Arg	Asp	Phe	Phe 365	Gly	Ser	Île	Ser	Gly 370	Ile	Pro	Leu	Pro	Pro 375
Pro	Pro	Asp	Leu	Leu 380	Pro	Lys	Met	Pro	Tyr 385	Glu	Pro	Leu	Thr	Pro 390
Val	Leu	Tyr	Leu	Ser 395	Leu	Trp	Asp	Ala	Leu 400	Lys	Tyr	Leu	Gly	Glu 405
Pro	Ile	Lys	Ser	Glu 410	Lys ·	Pro	Ile	Asn	Met 415	Glu	Asn	Leu	Pro	Val 420
Asn	Gly	Gly	Asn	Gly 425	Gln	Ser	Phe	Gly	Tyr 430	Ile	Leu	Tyr	Glu	Thr 435
Ser	Ile	Thr	Ser	Ser 440	Gly	Ile	Leu	Ser	Gly 445	His	Val	His	Asp	Arg 450
Gly	Gln	Val	Phe	Val 455	Asn	Thr	Val	Ser	Ile 460	Gly	Phe	Leu	Asp	Tyr 465
Lys	Thr	Thr	Lys	Ile 470	Ala	Val	Pro	Leu	Ile 475	Gln	Gly	Tyr	Thr	Val 480
Leu	Arg	Ile	Leu	Val 485	Glu	Asn	Arg	Gly	Arg 490	Val	Asn	Tyr	Gly	Glu 495
Asn	Ile	Asp	Asp	Gln	Arg	Lys	Gly	Leu	Ile	Gly	Asn	Leu	Tyr	Leu

	500	505	510
Asn Asp Ser Pr	b Leu Lys Asn	Phe Arg Ile Tyr	Ser Leu Asp Met
	515	520	525
Lys Lys Ser Ph	e Phe Gln Arg	Phe Gly Leu Asp	Lys Trp Xaa Ser
	530	535	540
Leu Pro Glu Th	r Pro Thr Leu 545	Pro Ala Phe Phe 550	Leu Gly Ser Leu 555
Ser Ile Ser Se	r Thr Pro Cys	Asp Thr Phe Leu	Lys Leu Glu Gly
	560	565	570
Trp Glu Lys Gl	y Val Val Phe	Ile Asn Gly Gln	Asn Leu Gly Arg
	575	580	585
Tyr Trp Asn Il	e Gly Pro Gln 590	Lys Thr Leu Tyr 595	Leu Pro Gly Pro 600
Trp Leu Ser Se	r Gly Ile Asn	Gln Val Ile Val	Phe Glu Glu Thr
	605	610	615
Met Ala Gly Pro	O Ala Leu Gln 620	Phe Thr Glu Thr 625	Pro His Leu Gly 630
Arg Asn Gln Ty	r Ile Lys 635		

<210> 176

<211> 2505

<212> DNA

<213> Homo Sapien

<400> 176

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gacettacat etgtgcagag tgggagatgg ggggtetece atcetggttg 650 cttcgaaaac ctgaaattca tctaagaacc tcagatccag acttccttgc 700 cgcagtggac tcctggttca aggtcttgct gcccaagata tatccatggc 750 tttatcacaa tgggggcaac atcattagca ttcaggtgga gaatgaatat 800 ggtagctaca gagcetgtga etteagetae atgaggeaet tggetggget 850 cttccgtgca ctgctaggag aaaagatctt gctcttcacc acagatgggc 900 ctgaaggact caagtgtggc tccctccggg gactctatac cactgtagat 950 tttggcccag ctgacaacat gaccaaaatc tttaccctgc ttcggaagta 1000 tgaaccccat gggccattgg taaactctga gtactacaca ggctggctgg 1050 attactgggg ccagaatcac tccacacggt ctgtgtcagc tgtaaccaaa 1100 ggactagaga acatgctcaa gttgggagcc agtgtgaaca tgtacatgtt 1150 ccatggaggt accaactttg gatattggaa tggtgccgat aagaagggac 1200 gcttccttcc gattactacc agctatgact atgatgcacc tatatctgaa 1250 gcaggggacc ccacacctaa gctttttgct cttcgagatg tcatcagcaa 1300 gttccaggaa gttcctttgg gacctttacc tcccccgagc cccaagatga 1350 tgcttggacc tgtgactctg cacctggttg ggcatttact ggctttccta 1400 gacttgcttt gcccccgtgg gcccattcat tcaatcttgc caatgacctt 1450 tgaggctgtc aagcaggacc atggcttcat gttgtaccga acctatatga 1500 cccataccat ttttgagcca acaccattct gggtgccaaa taatggagtc 1550 catgaccgtg cctatgtgat ggtggatggg gtgttccagg gtgttgtgga 1600 gcgaaatatg agagacaaac tatttttgac ggggaaactg gggtccaaac 1650 tggatatett ggtggagaac atggggagge teagetttgg gtetaacage 1700 agtgacttca agggcctgtt gaagccacca attctggggc aaacaatcct 1750 tacccagtgg atgatgttcc ctctgaaaat tgataacctt gtgaagtggt 1800 ggtttcccct ccagttgcca aaatggccat atcctcaagc tccttctggc 1850 cccacattct actccaaaac atttccaatt ttaggctcag ttggggacac 1900 atttctatat ctacctggat ggaccaaggg ccaagtctgg atcaatgggt 1950 ttaacttggg ccggtactgg acaaagcagg ggccacaaca gaccctctac 2000 gtgccaagat tcctgctgtt tcctagggga gccctcaaca aaattacatt 2050 gctggaacta gaagatgtac ctctccagcc ccaagtccaa tttttggata 2100 agectatect caatageact agtaetttge acaggacaca tateaattee 2150 cttteagetg atacactgag tgeetetgaa ceaatggagt taagtgggea 2200 ctgaaaggta ggeegggeat ggtggeteat geetgtaate eeageacttt 2250 gggaggetga gaegggtgga ttaeetgagg teaggaette aagaeeagee 2300 tggeeaacat ggtgaaacee egteteeact aaaaatacaa aaattageeg 2350 ggegtgatgg tgggeacete taateeeage taettgggag getgagggea 2400 ggagaattge ttgaateeag gaggeagagg ttgeagtgag tggaggttgt 2450 aceaetgeae teeageetgg etgaeagtga gaeaeteeat eteaaaaaa 2500 aaaaa 2505

<210> 177

<211> 654

<212> PRT

<213> Homo Sapien

<400> 177

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20 25 30

Val Val Asp Arg Gly His Asp Arg Phe Leu Leu Asp Gly Ala Pro
35 40 45

Phe Arg Tyr Val Ser Gly Ser Leu His Tyr Phe Arg Val Pro Arg
50 55 60

Val Leu Trp Ala Asp Arg Leu Leu Lys Met Arg Trp Ser Gly Leu 65 70 75

Asn Ala Ile Gln Phe Tyr Val Pro Trp Asn Tyr His Glu Pro Gln 80 85 90

Pro Gly Val Tyr Asn Phe Asn Gly Ser Arg Asp Leu Ile Ala Phe 95 100 105

Leu Asn Glu Ala Ala Leu Ala Asn Leu Leu Val Ile Leu Arg Pro
110 115 120

Gly Pro Tyr Ile Cys Ala Glu Trp Glu Met Gly Gly Leu Pro Ser 125 130 135

Trp Leu Leu Arg Lys Pro Glu Ile His Leu Arg Thr Ser Asp Pro 140 145 150

Asp Phe Leu Ala Ala Val Asp Ser Trp Phe Lys Val Leu Leu Pro 155 160 165

Lys Ile Tyr Pro Trp Leu Tyr His Asn Gly Gly Asn Ile Ile Ser 170 175 180

Ile	Gln	Val	Glu	Asn 185	Glu	Tyr	Gly	Ser	Tyr 190	Arg	Ala	Cys	Asp	Phe 195
Ser	Tyr	Met	Arg	His 200	Leu	Ala	Gly	Leu	Phe 205	Arg	Ala	Leu	Leu	Gly 210
Glu	Lys	Ile	Leu	Leu 215	Phe	Thr	Thr	Asp	Gly 220	Pro	Glu	Gly	Leu	Lys 225
Cys	Gly	Ser	Leu	Arg 230	_	Leu	Tyr	Thr	Thr 235	Val	Asp	Phe	Gly	Pro 240
Ala	Asp	Asn	Met	Thr 245	Lys	Ile	Phe	Thr	Leu 250	Leu	Arg	Lys	Tyr	Glu 255
Pro	His	Gly	Pro	Leu 260	Val	Asn	Ser	Glu	Tyr 265	Tyr	Thr	Gly	Trp	Leu 270
Asp	Tyr	Trp	Gly	Gln 275	Asn	His	Ser	Thr	Arg 280	Ser	Val	Ser	Ala	Val 285
Thr	Lys	Gly	Leu	Glu 290	Asn	Met	Leu	Lys	Leu 295	Gly	Ala	Ser	Val	Asn 300
Met	Tyr	Met	Phe	His 305	Gly	Gly	Thr	Asn	Phe 310	Gly	Tyr	Trp	Asn	Gly 315
Ala	Asp	Lys	Lys	Gly 320	Arg	Phe	Leu	Pro	Ile 325	Thr	Thr	Ser	Tyr	Asp 330
Tyr	Asp	Ala	Pro	Ile 335	Ser	Glu	Ala	Gly	Asp 340	Pro	Thr	Pro	Lys	Leu 345
Phe	Ala	Leu	Arg	Asp 350	Val	Ile	Ser	Lys	Phe 355	Gln	Glu	Val	Pro	Leu 360
Gly	Pro	Leu	Pro	Pro 365	Pro	Ser	Pro	Lys	Met 370	Met	Leu	Gly	Pro	Val 375
Thr	Leu	His	Leu	Val 380	Gly	His	Leu	Leu	Ala 385	Phe	Leu	Asp	Leu	Leu 390
Cys	Pro	Arg	Gly	Pro 395	Ile	His	Ser	Ile	Leu 400	Pro	Met	Thr	Phe	Glu 405
Ala	Val	Lys	Gln	Asp 410	His	Gly	Phe	Met	Leu 415	Tyr	Arg	Thr	Tyr	Met 420
Thr	His	Thr	Ile	Phe 425	Glu	Pro	Thr	Pro	Phe 430	Trp	Val	Pro	Asn	Asn 435
Gly	Val	His	Asp	Arg 440	Ala	Tyr	Val	Met	Val 445	Asp	Gly	Val	Phe	Gln 450
Gly	Val	Val	Glu	Arg 455	Asn	Met	Arg	Asp	Lys 460	Leu	Phe	Leu	Thr	Gly 465
Lys	Leu	Gly	Ser	Lys	Leu	Asp	Ile	Leu	Val	Glu	Asn	Met	Gly	Arg

				470					475					480
Leu	Ser	Phe	Gly	Ser 485	Asn	Ser	Ser	Asp	Phe 490	Lys	Gly	Leu	Leu	Lys 495
Pro	Pro	Ile	Leu	Gly 500	Gln	Thr	Ile	Leu	Thr 505	Gln	Trp	Met	Met	Phe 510
Pro	Leu	Lys	Ile	Asp 515	Asn	Leu	Val	Lys	Trp 520	Trp	Phe	Pro	Leu	Gln 525
Leu	Pro	Lys	Trp	Pro 530	Tyr	Pro	Gln	Ala	Pro 535	Ser	Gly	Pro	Thr	Phe 540
Tyr	Ser	Lys	Thr	Phe 545	Pro	Ile	Leu	Gly	Ser 550	Val	Gly	Asp	Thr	Phe 555
Leu	Tyr	Leu	Pro	Gly 560	Trp	Thr	Lys	Gly	Gln 565	Val	Trp	Ile	Asn	Gly 570
Phe	Asn	Leu	Gly	Arg 575	Tyr	Trp	Thr	Lys	Gln 580	Gly	Pro	Gln	Gln	Thr 585
Leu	Tyr	Val	Pro	Arg 590	Phe	Leu	Leu	Phe	Pro 595	Arg	Gly	Ala	Leu	Asn 600
Lvs	Ile	Thr	Leu	Leu	Glu	Leu	Glu	Asp	Val	Pro	Leu	Gln	Pro	Gln

Lys Ile Thr Leu Leu Glu Leu Glu Asp Val Pro Leu Gln Pro Gln 605 615

Val Gln Phe Leu Asp Lys Pro Ile Leu Asn Ser Thr Ser Thr Leu

620 625 630

His Arg Thr His Ile Asn Ser Leu Ser Ala Asp Thr Leu Ser Ala 635 640 645

Ser Glu Pro Met Glu Leu Ser Gly His 650

<210> 178

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 178

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<210> 179

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 179

tggacaaatc cccttgctca gccc 24

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<210> 180
<211> 50
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 180
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<210> 181
<211> 22
<212> DNA
<213> Artificial Sequence
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<400> 181
ccagctatga ctatgatgca cc 22
<210> 182
<211> 24
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<223> Synthetic Oligonucleotide Probe
<400> 182
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<210> 183
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<223> Synthetic Oligonucleotide Probe
<400> 183
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<210> 184
<211> 1947
<212> DNA
<213> Homo Sapien
<400> 184
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cagttacata tocattattt gtgtttatgg ctttatctgc ctctacactc 150
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- <210> 185
- <211> 501
- <212> PRT
- <213> Homo Sapien

<400> 185

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- Ile Cys Val Tyr Gly Phe Ile Cys Leu Tyr Thr Leu Phe Trp Leu 20 25 30
- Phe Arg Ile Pro Leu Lys Glu Tyr Ser Phe Glu Lys Val Arg Glu
 35 40 45
- Glu Ser Ser Phe Ser Asp Ile Pro Asp Val Lys Asn Asp Phe Ala
 50 55 60
- Phe Leu Leu His Met Val Asp Gln Tyr Asp Gln Leu Tyr Ser Lys
 65 70 75
- Arg Phe Gly Val Phe Leu Ser Glu Val Ser Glu Asn Lys Leu Arg 80 85 90
- Glu Ile Ser Leu Asn His Glu Trp Thr Phe Glu Lys Leu Arg Gln
 95 100 105
- His Ile Ser Arg Asn Ala Gln Asp Lys Gln Glu Leu His Leu Phe 110 115 120
- Met Leu Ser Gly Val Pro Asp Ala Val Phe Asp Leu Thr Asp Leu 125 130 135
- Asp Val Leu Lys Leu Glu Leu Ile Pro Glu Ala Lys Ile Pro Ala 140 145 150
- Lys Ile Ser Gln Met Thr Asn Leu Gln Glu Leu His Leu Cys His \$155\$
- Cys Pro Ala Lys Val Glu Gln Thr Ala Phe Ser Phe Leu Arg Asp 170 175 180
- His Leu Arg Cys Leu His Val Lys Phe Thr Asp Val Ala Glu Ile 185 190 195
- Pro Ala Trp Val Tyr Leu Leu Lys Asn Leu Arg Glu Leu Tyr Leu 200 205 210

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Ser	Leu	Arg	Glu	Leu 230	Arg	His	Leu	Lys	Ile 235	Leu	His	Val	Lys	Ser 240
Asn	Leu	Thr	Lys	Val 245	Pro	Ser	Asn	Ile	Thr 250	Asp	Val	Ala	Pro	His 255
Leu	Thr	Lys	Leu	Val 260	Ile	His	Asn	Asp	Gly 265	Thr	Lys	Leu	Leu	Val 270
Leu	Asn	Ser	Leu	Lys 275	Lys	Met	Met	Asn	Val 280	Ala	Glu	Leu	Glu	Leu 285
Gln	Asn	Cys	Glu	Leu 290	Glu	Arg	Ile	Pro	His 295	Ala	Ile	Phe	Ser	Leu 300
Ser	Asn	Leu	Gln	Glu 305	Leu	Asp	Leu	Lys	Ser 310	Asn	Asn	Ile	Arg	Thr 315
Ile	Glu	Glu	Ile	Ile 320	Ser	Phe	Gln	His	Leu 325	Lys	Arg	Leu	Thr	Cys 330
Leu	Lys	Leu	Trp	His 335	Asn	Lys	Ile	Val	Thr 340	Ile	Pro	Pro	Ser	Ile 345
Thr	His	Val	Lys	Asn 350		Glu	Ser	Leu	Tyr 355	Phe	Ser	Asn	Asn	Lys 360
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Ile Lys Arg	Gly Val Thr	Leu Asp Leu Gly	Leu Ile Asn Gln	Ser 380 Gln 395 Leu 410 Leu 425 Leu	Asn Pro Gly Ser	Leu Lys Gln Gln	Gln Gln Asn Leu	His Leu Cys	385 Leu 400 Phe 415 Ile 430 Gln 445	His Lys Thr Leu	Ile Cys Ser Glu	Thr Ile Leu Leu	Gly Lys Pro	Glu 390 Asn 405 Leu 420 Glu 435 Gly 450
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- Ser Cys Thr Val Ser Leu Gly Gly Ala Asn Met Ala Glu Thr His
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- Lys Ala Met Ile Leu Gln Leu Asn Pro Ser Glu Asn Cys Thr Trp
 50 55 60
- Thr Ile Glu Arg Pro Glu Asn Lys Ser Ile Arg Ile Ile Phe Ser
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Arg	Thr	Val	Phe	Val 140	Phe	Tyr	Tyr	Phe	Phe 145	Ser	Pro	Asn	Ile	Ser 150
Ile	Pro	Asn	Cys	Gly 155	Gly	Tyr	Leu	Asp	Thr 160	Leu	Glu	Gly	Ser	Phe 165
Thr	Ser	Pro	Asn	Tyr 170	Pro	Lys	Pro	His	Pro 175	Glu	Leu	Ala	Tyr	Cys 180
Val	Trp	His	Ile	Gln 185	Val	Glu	Lys	Asp	Tyr 190	Lys	Ile	Lys	Leu	Asn 195
Phe	Lys	Glu	Ile	Phe 200	Leu	Glu	Ile	Asp	Lys 205	Gln	Cys	Lys	Phe	Asp 210
Phe	Leu	Ala	Ile	Tyr 215	Asp	Gly	Pro	Ser	Thr 220	Asn	Ser	Gly	Leu	Ile 225
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Val	Glu	Phe	Ser	Val 320	Pro	Leu	Asn	Gly	Cys 325	Gly	Thr	Ile	Arg	Lys 330
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Ala	Ser	Ser	Thr	Ser 350	Glu	Val	Ile	Thr	Arg 355	Gln	Lys	Gln	Leu	Gln 360
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- <212> PRT
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- Leu Pro Ala Trp Phe Asp Gln Ala Lys Phe Gly Ile Phe Ile His
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- Trp Gly Val Phe Ser Val Pro Ser Phe Gly Ser Glu Trp Phe Trp
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- Trp Tyr Trp Gln Lys Glu Lys Ile Pro Lys Tyr Val Glu Phe Met
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- Lys Asp Asn Tyr Pro Pro Ser Phe Lys Tyr Glu Asp Phe Gly Pro 95 100 105
- Leu Phe Thr Ala Lys Phe Phe Asn Ala Asn Gln Trp Ala Asp Ile 110 115 120
- Phe Gln Ala Ser Gly Ala Lys Tyr Ile Val Leu Thr Ser Lys His
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- His Glu Gly Phe Thr Leu Trp Gly Ser Glu Tyr Ser Trp Asn Trp
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- Asn Ala Ile Asp Glu Gly Pro Lys Arg Asp Ile Val Lys Glu Leu 155 : 160 165
- Glu Val Ala Ile Arg Asn Arg Thr Asp Leu Arg Phe Gly Leu Tyr 170 175 180
- Tyr Ser Leu Phe Glu Trp Phe His Pro Leu Phe Leu Glu Asp Glu 185 190 195
- Ser Ser Ser Phe His Lys Arg Gln Phe Pro Val Ser Lys Thr Leu 200 205 210
- Pro Glu Leu Tyr Glu Leu Val Asn Asn Tyr Gln Pro Glu Val Leu 215 220 225
- Trp Ser Asp Gly Asp Gly Gly Ala Pro Asp Gln Tyr Trp Asn Ser 230 235 240

Thr Gly Phe Leu Ala Trp Leu Tyr Asn Glu Ser Pro Val Arg Gly Thr Val Val Thr Asn Asp Arg Trp Gly Ala Gly Ser Ile Cys Lys 260 265 His Gly Gly Phe Tyr Thr Cys Ser Asp Arg Tyr Asn Pro Gly His Leu Leu Pro His Lys Trp Glu Asn Cys Met Thr Ile Asp Lys Leu Ser Trp Gly Tyr Arg Arg Glu Ala Gly Ile Ser Asp Tyr Leu Thr Ile Glu Glu Leu Val Lys Gln Leu Val Glu Thr Val Ser Cys Gly Gly Asn Leu Leu Met Asn Ile Gly Pro Thr Leu Asp Gly Thr Ile Ser Val Val Phe Glu Glu Arg Leu Arg Gln Val Gly Ser Trp Leu Lys Val Asn Gly Glu Ala Ile Tyr Glu Thr Tyr Thr Trp Arg Ser Gln Asn Asp Thr Val Thr Pro Asp Val Trp Tyr Thr Ser Lys Pro Lys Glu Lys Leu Val Tyr Ala Ile Phe Leu Lys Trp Pro Thr Ser 400 Gly Gln Leu Phe Leu Gly His Pro Lys Ala Ile Leu Gly Ala Thr Glu Val Lys Leu Leu Gly His Gly Gln Pro Leu Asn Trp Ile Ser 425 Leu Glu Gln Asn Gly Ile Met Val Glu Leu Pro Gln Leu Thr Ile His Gln Met Pro Cys Lys Trp Gly Trp Ala Leu Ala Leu Thr Asn 455 460

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Ser Ile Gln Val Ser Cys Arg Ile Met Gly Ile Thr Leu $^{\prime}$ Val Ser 35 40 45

Lys Lys Ala Asn Gln Gln Leu Asn Phe Thr Glu Ala Lys Glu Ala 50 55 60

Cys Arg Leu Leu Gly Leu Ser Leu Ala Gly Lys Asp Gln Val Glu
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Thr Ala Leu Lys Ala Ser Phe Glu Thr Cys Ser Tyr Gly Trp Val 80 85 90

Gly Asp Gly Phe Val Val Ile Ser Arg Ile Ser Pro Asn Pro Lys 95 100 105

Cys Gly Lys Asn Gly Val Gly Val Leu Ile Trp Lys Val Pro Val
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Ser Arg Gln Phe Ala Ala Tyr Cys Tyr Asn Ser Ser Asp Thr Trp
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Thr Asn Ser Cys Ile Pro Glu Ile Ile Thr Thr Lys Asp Pro Ile 140 145 150

Phe Asn Thr Gln Thr Ala Thr Gln Thr Thr Glu Phe Ile Val Ser
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Asp Ser Thr Tyr Ser Val Ala Ser Pro Tyr Ser Thr Ile Pro Ala 170 175 180

Pro Thr Thr Pro Pro Ala Pro Ala Ser Thr Ser Ile Pro Arg
185 190 195

Arg Lys Lys Leu Ile Cys Val Thr Glu Val Phe Met Glu Thr Ser 200 205 Thr Met Ser Thr Glu Thr Glu Pro Phe Val Glu Asn Lys Ala Ala Phe Lys Asn Glu Ala Ala Gly Phe Gly Gly Val Pro Thr Ala Leu Leu Val Leu Ala Leu Leu Phe Phe Gly Ala Ala Ala Gly Leu Gly Phe Cys Tyr Val Lys Arg Tyr Val Lys Ala Phe Pro Phe Thr Asn 260 Lys Asn Gln Gln Lys Glu Met Ile Glu Thr Lys Val Val Lys Glu 275 280 Glu Lys Ala Asn Asp Ser Asn Pro Asn Glu Glu Ser Lys Lys Thr Asp Lys Asn Pro Glu Glu Ser Lys Ser Pro Ser Lys Thr Thr Val Arg Cys Leu Glu Ala Glu Val <210> 202 <211> 24 <212> DNA <213> Artificial Sequence <223> Synthetic Oligonucleotide Probe <400> 202 gagctttcca tccaggtgtc atgc 24 <210> 203 <211> 22 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 203 gtcagtgaca gtacctactc gg 22 <210> 204 <211> 24 <212> DNA <213> Artificial Sequence <223> Synthetic Oligonucleotide Probe <400> 204

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- <223> unknown base
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35 40 45

Cys His Gly Leu Pro Thr Gln Arg Glu Asp Gly Asn Pro Cys Asp
50 55 60

Phe Asp Trp Arg Glu Val Glu Ile Leu Met Phe Leu Ser Ala Ile
65 70 75

Val Met Met Lys Asn Arg Arg Ser Ile Thr Val Glu Gln His Ile 80 85 90

Gly Asn Ile Phe Met Phe Ser Lys Val Ala Asn Thr Ile Leu Phe 95 100 105

Phe Arg Leu Asp Ile Arg Met Gly Leu Leu Tyr Ile Thr Leu Cys
110 115 120

Ile Val Phe Leu Met Thr Cys Lys Pro Pro Leu Tyr Met Gly Pro 130 Glu Tyr Ile Lys Tyr Phe Asn Asp Lys Thr Ile Asp Glu Glu Leu Glu Arg Asp Lys Arg Val Thr Trp Ile Val Glu Phe Phe Ala Asn Trp Ser Asn Asp Cys Gln Ser Phe Ala Pro Ile Tyr Ala Asp Leu 170 175 Ser Leu Lys Tyr Asn Cys Thr Gly Leu Asn Phe Gly Lys Val Asp 190 Val Gly Arg Tyr Thr Asp Val Ser Thr Arg Tyr Lys Val Ser Thr 205 Ser Pro Leu Thr Lys Gln Leu Pro Thr Leu Ile Leu Phe Gln Gly 215 220 225 Gly Lys Glu Ala Met Arg Arg Pro Gln Ile Asp Lys Lys Gly Arg 230 235 Ala Val Ser Trp Thr Phe Ser Glu Glu Asn Val Ile Arg Glu Phe 245 Asn Leu Asn Glu Leu Tyr Gln Arg Ala Lys Lys Leu Ser Lys Ala 260 265 Gly Asp Asn Ile Pro Glu Glu Gln Pro Val Ala Ser Thr Pro Thr 285 275 280 Thr Val Ser Asp Gly Glu Asn Lys Lys Asp Lys 290 295 <210> 208 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 208 gcttggatat tcgcatgggc ctac 24 <210> 209 <211> 20 <212> DNA <213> Artificial Sequence <223> Synthetic Oligonucleotide Probe <400> 209 tggagacaat atccctgagg 20 <210> 210

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Gly	Val	Lys	Leu	Val 50	Val	Glu	Thr	Pro	Glu 55	Glu	Thr	Leu	Phe	Thr 60
Tyr	Gln	Gly	Ala	Ser 65	Val	Ile	Leu	Pro	Cys 70	Arg	Tyr	Arg	Tyr	Glu 75
Pro	Ala	Leu	Val	Ser 80	Pro	Arg	Arg	Val	Arg 85	Val	Lys	Trp	Trp	Lys 90
Leu	Ser	Glu	Asn	Gly 95	Ala	Pro	Glu	Lys	Asp 100	Val	Leu	Val	Ala	Ile 105
Gly	Leu	Arg	His	Arg 110	Ser	Phe	Gly	Asp	Tyr 115	Gln	Gly	Arg	Val	His 120
Leu	Arg	Gln	Asp	Lys 125	Glu	His	Asp	Val	Ser 130	Leu	Glu	Ile	Gln	Asp 135
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Val	Val	Phe	Pro	Tyr 170	Gln	Ser	Pro	Asn	Gly 175	Arg	Tyr	Gln	Phe	Asn 180
Phe	His	Glu	Gly		Gln	Val	Cys	Ala	_	Gln	Ala	Ala	Val	
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Trp	Cys	Asn	Ala	Gly 215	Trp	Leu	Gln	Asp	Ala 220	Thr	Val	Gln	Tyr	Pro 225
Ile	Met	Leu	Pro	Arg 230	Gln	Pro	Cys	Gly	Gly 235	Pro	Gly	Leu	Ala	Pro 240
Gly	Val	Arg	Ser	Tyr 245	Gly	Pro	Arg	His	Arg 250	Arg	Leu	His	Arg	Tyr 255
Asp	Val	Phe	Cys	Phe 260	Ala	Thr	Ala	Leu	Lys 265	Gly	Arg	Val	Tyr	Tyr 270
Leu	Glu	His	Pro	Glu 275	Lys	Leu	Thr	Leu	Thr 280	Glu	Ala	Arg	Glu	Ala 285
Cys	Gln	Glu	Asp	Asp 290	Ala	Thr	Ile	Ala	Lys 295	Val	Gly	Ģln	Leu	Phe 300
Ala	Ala	Trp	Lys	Phe 305	His	Gly	Leu	Asp	Arg 310	Cys	Asp	Ala	Gly	Trp 315
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Asp Asp Ala His Gly Asn Phe Gln Tyr Asp His Glu Ala Phe Leu
50 55 60

Gly Arg Glu Val Ala Lys Glu Phe Asp Gln Leu Thr Pro Glu Glu
65 70 75

Ser Gln Ala Arg Leu Gly Arg Ile Val Asp Arg Met Asp Arg Ala 80 85 90

Gly Asp Gly Asp Gly Trp Val Ser Leu Ala Glu Leu Arg Ala Trp 95 100 105

Ile Ala His Thr Gln Gln Arg His Ile Arg Asp Ser Val Ser Ala 110 115 120

Ala Trp Asp Thr Tyr Asp Thr Asp Arg Asp Gly Arg Val Gly Trp
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Glu Glu Leu Arg Asn Ala Thr Tyr Gly His Tyr Ala Pro Gly Glu 140 145 150

Glu Phe His Asp Val Glu Asp Ala Glu Thr Tyr Lys Lys Met Leu

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Ala Ile Thr Cys Phe Thr Arg Gly Leu Asp Ile Arg Lys Glu Lys
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Ala Asp Val Leu Cys Pro Gly Gly Cys Pro Leu Glu Glu Phe Ser
50 55 60

Val Tyr Gly Asn Ile Val Tyr Ala Ser Val Ser Ser Ile Cys Gly 65 70 75

Ala Ala Val His Arg Gly Val Ile Ser Asn Ser Gly Gly Pro Val 80 85 90

Arg Val Tyr Ser Leu Pro Gly Arg Glu Asn Tyr Ser Ser Val Asp 95 100 105

Ala Asn Gly Ile Gln Ser Gln Met Leu Ser Arg Trp Ser Ala Ser 110 115 120

Phe Thr Val Thr Lys Gly Lys Ser Ser Thr Gln Glu Ala Thr Gly
125 130 135

Gln Ala Val Ser Thr Ala His Pro Pro Thr Gly Lys Arg Leu Lys 140 145 150

Lys Thr Pro Glu Lys Lys Thr Gly Asn Lys Asp Cys Lys Ala Asp 155 160 165

Ile Ala Phe Leu Ile Asp Gly Ser Phe Asn Ile Gly Gln Arg Arg 170 175 180

Phe Asn Leu Gln Lys Asn Phe Val Gly Lys Val Ala Leu Met Leu 185 190 195

Gly Ile Gly Thr Glu Gly Pro His Val Gly Leu Val Gln Ala Ser 200 205 210

Glu His Pro Lys Ile Glu Phe Tyr Leu Lys Asn Phe Thr Ser Ala 215 220 225

Lys Asp Val Leu Phe Ala Ile Lys Glu Val Gly Phe Arg Gly Gly 230 235 240

Asn Ser Asn Thr Gly Lys Ala Leu Lys His Thr Ala Gln Lys Phe 245 250 255

Phe	Thr	Val	Asp	Ala 260	Gly	Val	Arg	Lys	Gly 265	Ile	Pro	Lys	Val	Val 270
Val	Val	Phe	Ile	Asp 275	Gly	Trp	Pro	Ser	Asp 280	Asp	Ile	Glu	Glu	Ala 285
Gly	Ile	Val	Ala	Arg 290	Glu	Phe	Gly	Val	Asn 295	Val	Phe	Ile	Val	Ser 300
Val	Ala	Lys	Pro	Ile 305	Pro	Glu	Glu	Leu	Gly 310	Met	Val	Gln	Asp	Val 315
Thr	Phe	Val	Asp	Lys 320	Ala	Val	Cys	Arg	Asn 325	Asn	Gly	Phe	Phe	Ser 330
Tyr	His	Met	Pro	Asn 335	Trp	Phe	Gly	Thr	Thr 340	Lys	Tyr	Val	Lys	Pro 345
Leu	Val	Gln	Lys	Leu 350	Cys	Thr	His	Glu	Gln 355	Met	Met	Cys	Ser	Lys 360
Thr	Cys	Tyr	Asn	Ser 365	Val	Asn	Ile	Ala	Phe 370	Leu	Ile	Asp	Gly	Ser 375
Ser	Ser	Val	Gly	Asp 380	Ser	Asn	Phe	Arg	Leu 385	Met	Leu	Glu	Phe	Val 390
Ser	Asn	Ile	Ala	Lys 395	Thr	Phe	Glu	Ile	Ser 400	Asp	Ile	Gly	Ala	Lys 405
Ile	Ala	Ala	Val	Gln 410	Phe	Thr	Tyr	Asp	Gln 415	Arg	Thr	Glu	Phe	Ser 420
Phe	Thr	Asp	Tyr	Ser 425	Thr	Lys	Glu	Asn	Val 430	Leu	Ala	Val	Ile	Arg 435
Asn	Ile	Arg	Tyr	Met 440	Ser	Gly	Gly	Thr	Ala 445	Thr	Gly	Asp	Ala	Ile 450
Ser	Phe	Thr	Val	Arg 455	Asn	Val	Phe	Gly	Pro 460	Ile	Arg	Glu	Ser	Pro 465
Asn	Lys	Asn	Phe	Leu 470	Val	Ile	Val	Thr	Asp 475	Gly	Gln	Ser	Tyr	Asp 480
Asp	Val	Gln	Gly	Pro 485	Ala	Ala	Ala	Ala	His 490	Asp	Ala	Gly	Ile	Thr 495
Ile	Phe	Ser	Val	Gly 500	Val	Ala	Trp	Ala	Pro 505	Leu	Asp	Asp	Leu	Lys 510
Asp	Met	Ala	Ser	Lys 515	Pro	Lys	Glu	Ser	His 520	Ala	Phe	Phe	Thr	Arg 525
Glu	Phe	Thr	Gly	Leu 530	Glu	Pro	Ile	Val	Ser 535	Asp	Val	Ile	Arg	Gly 540
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Pro Val Lys Pro Gly Pro Ala Leu Ser Tyr Pro Gln Glu Glu Ala 35 40 45

Thr Leu Asn Glu Met Phe Arg Glu Val Glu Glu Leu Met Glu Asp
50 55 60

Thr Gln His Lys Leu Arg Ser Ala Val Glu Glu Met Glu Ala Glu
65 70 75

Glu Ala Ala Ala Lys Ala Ser Ser Glu Val Asn Leu Ala Asn Leu 80 85 90

Pro Pro Ser Tyr His Asn Glu Thr Asn Thr Asp Thr Lys Val Gly
95 100 105

Asn Asn Thr Ile His Val His Arg Glu Ile His Lys Ile Thr Asn 110 . 115 120

Asn Gln Thr Gly Gln Met Val Phe Ser Glu Thr Val Ile Thr Ser 125 130 135

Val Gly Asp Glu Glu Gly Arg Arg Ser His Glu Cys Ile Ile Asp 140 145 150

Glu Asp Cys Gly Pro Ser Met Tyr Cys Gln Phe Ala Ser Phe Gln 155 160 165

Tyr Thr Cys Gln Pro Cys Arg Gly Gln Arg Met Leu Cys Thr Arg
170 175 180

Asp Ser Glu Cys Cys Gly Asp Gln Leu Cys Val Trp Gly His Cys 185 190 195

Thr Lys Met Ala Thr Arg Gly Ser Asn Gly Thr Ile Cys Asp Asn 200 205 210

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Gln Arg Asp Cys Gln Pro Gly Leu Cys Cys Ala Phe Gln Arg Gly
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 Cys His Asp Pro Ala Ser Arg Leu Leu Asp Leu Ile Thr Trp Glu
Leu Glu Pro Asp Gly Ala Leu Asp Arg Cys Pro Cys Ala Ser Gly
Leu Leu Cys Gln Pro His Ser His Ser Leu Val Tyr Val Cys Lys
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 Pro Thr Phe Val Gly Ser Arg Asp Gln Asp Gly Glu Ile Leu Leu
 Pro Arg Glu Val Pro Asp Glu Tyr Glu Val Gly Ser Phe Met Glu
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Dro Cla Cys Ala Cys Cla Lla Arg Dro Trp Thy The Dro Arg Com

Pro Gln Cys Ala Cys Gln Ile Arg Pro Trp Tyr Thr Pro Arg Ser 35 40 45

Ser Tyr Arg Glu Ala Thr Thr Val Asp Cys Asn Asp Leu Phe Leu
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Thr Ala Val Pro Pro Ala Leu Pro Ala Gly Thr Gln Thr Leu Leu 65 70 75

Leu Gln Ser Asn Ser Ile Val Arg Val Asp Gln Ser Glu Leu Gly
80 85 90

Tyr Leu Ala Asn Leu Thr Glu Leu Asp Leu Ser Gln Asn Ser Phe 95 100 105

Ser Asp Ala Arg Asp Cys Asp Phe His Ala Leu Pro Gln Leu Leu 110 115 120

Ser Leu His Leu Glu Glu Asn Gln Leu Thr Arg Leu Glu Asp His 125 130 130

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His	Leu	Lys	Glu	Leu 290	Gly	Leu	Asn	Asn	Met 295	Glu	Glu	Leu	Val	Ser 300
Ile	Asp	Lys	Phe	Ala 305	Leu	Val	Asn	Leu	Pro 310	Glu	Leu	Thr	Lys	Leu 315
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Phe	His	His	Leu	Pro 335	Gln	Met	Glu	Thr	Leu 340	Met	Leu	Asn	Asn	Asn 345
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Val ;	Ile	Arg	Trp	Ala 380	Asn	Ala	Thr	Gly	Thr 385	Arg	Val	Arg	Phe	Ile 390
Glu	Pro	Gln	Ser	Thr 395	Leu	Cys	Ala	Glu	Pro 400	Pro	Asp	Leu	Gln	Arg 405
Leu	Pro	Val	Arg	Glu 410	Val	Pro	Phe	Arg	Glu 415	Met	Thr	Asp	His	Cys 420
Leu	Pro	Leu	Ile	Ser 425	Pro	Arg	Ser	Phe	Pro 430	Pro	Ser	Leu	Gln	Val 435
Ala	Ser	Gly	Glu	Ser 440	Met	Val	Leu	His	Cys 445	Arg	Ala	Leu	Ala	Glu 450

Pro	Glu	Pro	Glu	Ile 455	Tyr	Trp	Val	Thr	Pro 460	Ala	Gly	Leu	Arg	Leu 465
Thr	Pro	Ala	His	Ala 470	Gly	Arg	Arg	Tyr	Arg 475	Val	Tyr	Pro	Glu	Gly 480
Thr	Leu	Glu	Leu	Arg 485	Arg	Val	Thr	Ala	Glu 490	Glu	Ala	Gly	Leu	Tyr 495
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Ser	Val	Val	Val	Gly 515	Arg	Ala	Leu	Leu	Gln 520	Pro	Gly	Arg	Asp	Glu 525
Gly	Gln	Gly	Leu	Glu 530	Leu	Arg	Val	Gln	Glu 535	Thr	His	Pro	Tyr	His 540
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Leu	Thr	Trp	Ser	Ser 560	Ala	Ser	Ser	Leu	Arg 565	Gly	Gln	Gly	Ala	Thr 570
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Arg	Leu	Leu	Gln	Ala 590	Thr	Glu	Tyr	Trp	Ala 595	Cys	Leu	Gln	Val	Ala 600
Phe	Ala	Asp	Ala	His 605	Thr	Gln	Leu	Ala	Cys 610	Val	Trp	Ala	Arg	Thr 615
Lys	Glu	Ala	Thr	Ser 620	Cys	His	Arg	Ala	Leu 625	Gly	Asp	Arg	Pro	Gly 630
Leu	Ile	Ala	Ile	Leu 635	Ala	Leu	Ala	Val	Leu 640	Leu	Leu	Ala	Ala	Gly 645
Leu	Ala	Ala	His	Leu 650	Gly	Thr	Gly	Gln	Pro 655	Arg	Lys	Gly	Val	Gly 660
Gly	Arg	Arg	Pro	Leu 665	Pro	Pro	Ala	Trp	Ala 670	Phe	Trp	Gly	Trp	Ser 675
Ala	Pro	Ser	Val	Arg 680	Val	Val	Ser	Ala	Pro 685	Leu	Val	Leu	Pro	Trp 690
Asn	Pro	Gly	Arg	Lys 695	Leu	Pro	Arg	Ser	Ser 700		Gly	Glu	Thr	Leu .705
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<211> 546

<212> PRT

<213> Homo Sapien

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Cys	Thr	Val	Asp	Ile 35	Glu	Ser	Leu	Thr	Gly 40	Tyr	Arg	Thr	Tyr	Arg 45
Cys	Ala	His	Pro	Leu 50	Ala	Thr	Leu	Phe	Lys 55	Ile	Leu	Ala	Ser	Phe 60
Tyr	Ile	Ser	Leu	Val 65	Ile	Phe	Tyr	Gly	Leu 70	Ile	Cys	Met	Tyr	Thr 75
Leu	Trp	Trp	Met	Leu 80	Arg	Arg	Ser	Leu	Lys 85	Lys	Tyr	Ser	Phe	Glu 90
Ser	Ile	Arg	Glu	Glu 95	Ser	Ser	Tyr	Ser	Asp 100	Ile	Pro	Asp	Val	Lys 105
Asn	Asp	Phe	Ala	Phe 110	Met	Leu	His	Leu	Ile 115	Asp	Gln	Tyr	Asp	Pro 120
Leu	Tyr	Ser	Lys	Arg 125	Phe	Ala	Val	Phe	Leu 130	Ser	Glu	Val	Ser	Glu 135
Asn	Lys	Leu	Arg	Gln 140	Leu	Asn	Leu	Asn	Asn 145	Glu	Trp	Thr	Leu	Asp 150
Lys	Leu	Arg	Gln	Arg 155	Leu	Thr	Lys	Asn	Ala 160	Gln	Asp	Lys	Leu	Glu 165
Leu	His	Leu	Phe	Met 170	Leu	Ser	Gly	Ile	Pro 175	Asp	Thr	Val	Phe	Asp 180
Leu	Val	Glu	Leu	Glu 185	Val	Leu	Lys	Leu	Glu 190	Leu	Ile	Pro	Asp	Val 195
Thr	Ile	Pro	Pro	Ser 200	Ile	Ala	Gln	Leu	Thr 205	Gly	Leu	Lys	Glu	Leu 210
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	Leu	Tyr	His	Thr 215	Ala	Ala	Lys	Ile		Ala	Pro	Ala	Leu	Ala 225
Phe				215					Glu 220		Pro Lys			225
	Leu	Arg	Glu	215 Asn 230	Leu	Arg	Ala	Leu	Glu 220 His 235	Ile		Phe	Thr	225 Asp 240
Ile	Leu Lys	Arg Glu	Glu Ile	215 Asn 230 Pro 245	Leu Leu	Arg Trp	Ala Ile	Leu Tyr	Glu 220 His 235 Ser 250	Ile Leu	Lys	Phe Thr	Thr Leu	225 Asp 240 Glu 255
Ile Glu	Leu Lys Leu	Arg Glu His	Glu Ile Leu	215 Asn 230 Pro 245 Thr 260	Leu Leu Gly	Arg Trp Asn	Ala Ile Leu	Leu Tyr Ser	Glu 220 His 235 Ser 250 Ala 265	Ile Leu Glu	Lys Lys	Phe Thr Asn	Thr Leu Arg	225 Asp 240 Glu 255 Tyr

Val Gly Val His Leu Gln Lys Leu Ser Ile Asn Asn Glu Gly Thr 305 310 Lys Leu Ile Val Leu Asn Ser Leu Lys Lys Met Ala Asn Leu Thr Glu Leu Glu Leu Ile Arg Cys Asp Leu Glu Arg Ile Pro His Ser Ile Phe Ser Leu His Asn Leu Gln Glu Ile Asp Leu Lys Asp Asn Asn Leu Lys Thr Ile Glu Glu Ile Ile Ser Phe Gln His Leu His 370 Arg Leu Thr Cys Leu Lys Leu Trp Tyr Asn His Ile Ala Tyr Ile Pro Ile Gln Ile Gly Asn Leu Thr Asn Leu Glu Arg Leu Tyr Leu Asn Arg Asn Lys Ile Glu Lys Ile Pro Thr Gln Leu Phe Tyr Cys 415 Arg Lys Leu Arg Tyr Leu Asp Leu Ser His Asn Asn Leu Thr Phe 430 Leu Pro Ala Asp Ile Gly Leu Leu Gln Asn Leu Gln Asn Leu Ala 440 445 Ile Thr Ala Asn Arg Ile Glu Thr Leu Pro Pro Glu Leu Phe Gln 455 460 Cys Arg Lys Leu Arg Ala Leu His Leu Gly Asn Asn Val Leu Gln 470 475 Ser Leu Pro Ser Arg Val Gly Glu Leu Thr Asn Leu Thr Gln Ile 485 490 Glu Leu Arg Gly Asn Arg Leu Glu Cys Leu Pro Val Glu Leu Gly 500 505 Glu Cys Pro Leu Leu Lys Arg Ser Gly Leu Val Val Glu Glu Asp 515 Leu Phe Asn Thr Leu Pro Pro Glu Val Lys Glu Arg Leu Trp Arg 530 Ala Asp Lys Glu Gln Ala

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<213> Artificial Sequence

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<223> Synthetic Oligonucleotide Probe

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Arg Lys Asp Ala Tyr Met Phe Trp Trp Leu Tyr Tyr Ala Thr Asn
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<210> 255

<211> 452

<212> PRT

<213> Homo Sapien

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Gly Phe Ser	Tyr Val 125	Asn	Gly	Ser	Gly	Ala 130	Tyr	Ala	Lys	Asp	Leu 135
Ala Met Val	Ala Ser 140	Asp	Met	Met	Val	Leu 145	Leu	Lys	Thr	Phe	Phe 150
Ser Cys His	Lys Glu 155	Phe	Gln	Thr	Val	Pro 160	Phe	Tyr	Ile	Phe	Ser 165
Glu Ser Tyr	Gly Gly 170	Lys	Met	Ala	Ala	Gly 175	Ile	Gly	Leu	Glu	Leu 180
Tyr Lys Ala	Ile Gln 185	Arg	Gly	Thr	Ile	Lys 190	Cys	Asn	Phe	Ala	Gly 195
Val Ala Leu	Gly Asp 200	Ser	Trp	Ile	Ser	Pro 205	Val	Asp	Ser	Val	Leu 210
Ser Trp Gly	Pro Tyr 215	Leu	Tyr	Ser	Met	Ser 220	Leu	Leu	Glu	Asp	Lys 225
Gly Leu Ala	Glu Val 230	Ser	Lys	Val	Ala	Glu 235	Gln	Val	Leu	Asn	Ala 240
Val Asn Lys	Gly Leu 245	Tyr	Arg	Glu	Ala	Thr 250	Glu	Leu	Trp	Gly	Lys 255
Ala Glu Met	Ile Ile 260	Glu	Gln	Asn	Thr	Asp 265	Gly	Val	Asn	Phe	Tyr 270
Asn Ile Leu	Thr Lys 275	Ser	Thr	Pro	Thr	Ser 280	Thr	Met	Glu	Ser	Ser 285
Leu Glu Phe	Thr Gln 290	Ser	His	Leu	Val	Cys 295	Leu	Cys	Gln	Arg	His 300
Val Arg His	Leu Gln 305	Arg	Asp	Ala	Leu	Ser 310	Gln	Leu	Met	Asn	Gly 315
Pro Ile Arg	Lys Lys	Leu	Lys	Ile	Ile	Pro 325	Glu	Asp	Gln	Ser	Trp 330
Gly Gly Gln	Ala Thr 335	Asn	Val	Phe	Val	Asn 340	Met	Glu	Glu	Asp	Phe 345
Met Lys Pro	Val Ile 350	Ser	Ile	Val	Asp	Glu 355	Leu	Leu	Gļu	Ala	Gly 360
Ile Asn Val	Thr Val	Tyr .	Asn	Gly	Gln	Leu 370	Asp	Leu	Ile	Val	Asp 375

Thr Met Gly Gln Glu Ala Trp Val Arg Lys Leu Lys Trp Pro Glu 380 385 390

Leu Pro Lys Phe Ser Gln Leu Lys Trp Lys Ala Leu Tyr Ser Asp 395 400 405

Pro Lys Ser Leu Glu Thr Ser Ala Phe Val Lys Ser Tyr Lys Asn 410 415 420

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Gln Glu

<210> 256

<211> 1100

<212> DNA

<213> Homo Sapien

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- <211> 314
- <212> PRT
- <213> Homo Sapien

<400> 257

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- Gly Pro Cys Gly Arg Arg Val Ile Thr Ser Arg Ile Val Gly Gly
 35 40 45
- Glu Asp Ala Glu Leu Gly Arg Trp Pro Trp Gln Gly Ser Leu Arg
 50 55 60
- Leu Trp Asp Ser His Val Cys Gly Val Ser Leu Leu Ser His Arg
 65 70 75
- Trp Ala Leu Thr Ala Ala His Cys Phe Glu Thr Tyr Ser Asp Leu 80 85 90
- Ser Asp Pro Ser Gly Trp Met Val Gln Phe Gly Gln Leu Thr Ser 95 100 105
- Met Pro Ser Phe Trp Ser Leu Gln Ala Tyr Tyr Thr Arg Tyr Phe
 110 115 120
- Val Ser Asn Ile Tyr Leu Ser Pro Arg Tyr Leu Gly Asn Ser Pro 125 130 135
- Tyr Asp Ile Ala Leu Val Lys Leu Ser Ala Pro Val Thr Tyr Thr 140 145 150
- Lys His Ile Gln Pro Ile Cys Leu Gln Ala Ser Thr Phe Glu Phe
 155 160 165
- Glu Asn Arg Thr Asp Cys Trp Val Thr Gly Trp Gly Tyr Ile Lys
 170 175 180
- Glu Asp Glu Ala Leu Pro Ser Pro His Thr Leu Gln Glu Val Gln
 185 190 195
- Val Ala Ile Ile Asn Asn Ser Met Cys Asn His Leu Phe Leu Lys
 200 205 210
- Tyr Ser Phe Arg Lys Asp Ile Phe Gly Asp Met Val Cys Ala Gly 215 220 225

Asn Ala Gln Gly Gly Lys Asp Ala Cys Phe Gly Asp Ser Gly Gly 240

Pro Leu Ala Cys Asn Lys Asn Gly Leu Trp Tyr Gln Ile Gly Val 255

Val Ser Trp Gly Val Gly Cys Gly Arg Pro Asn Arg Pro Gly Val 270

Tyr Thr Asn Ile Ser His His Phe Glu Trp 280 Ile Gln Lys Leu Met 285

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<400> 258

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<210> 258

<211> 2427

<212> DNA

<213> Homo Sapien

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- <210> 259
- <211> 556
- <212> PRT
- <213> Homo Sapien

<400> 259

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- Leu Pro Pro Gly Trp Val Ser Leu Gly Arg Ala Asp Pro Glu Glu
 45
- Glu Leu Ser Leu Thr Phe Ala Leu Arg Gln Gln Asn Val Glu Arg
 50 55 60
- Leu Ser Glu Leu Val Gln Ala Val Ser Asp Pro Ser Ser Pro Gln 65 70 75
- Tyr Gly Lys Tyr Leu Thr Leu Glu Asn Val Ala Asp Leu Val Arg 80 85 90
- Pro Ser Pro Leu Thr Leu His Thr Val Gln Lys Trp Leu Leu Ala 95 100 105
- Ala Gly Ala Gln Lys Cys His Ser Val Ile Thr Gln Asp Phe Leu 110 115 120
- Thr Cys Trp Leu Ser Ile Arg Gln Ala Glu Leu Leu Pro Gly
 125 130 135
- Ala Glu Phe His His Tyr Val Gly Gly Pro Thr Glu Thr His Val 140 145 150
- Val Arg Ser Pro His Pro Tyr Gln Leu Pro Gln Ala Leu Ala Pro 155 160 165
- His Val Asp Phe Val Gly Gly Leu His Arg Phe Pro Pro Thr Ser 170 175 180
- Ser Leu Arg Gln Arg Pro Glu Pro Gln Val Thr Gly Thr Val Gly
 185 190 190
- Leu His Leu Gly Val Thr Pro Ser Val Ile Arg Lys Arg Tyr Asn 200 205 210
- Leu Thr Ser Gln Asp Val Gly Ser Gly Thr Ser Asn Asn Ser Gln
 215
 220
 225
- Ala Cys Ala Gln Phe Leu Glu Gln Tyr Phe His Asp Ser Asp Leu 230 235 240
- Ala Gln Phe Met Arg Leu Phe Gly Gly Asn Phe Ala His Gln Ala

				245					250					255
Ser	Val	Ala	Arg	Val 260	Val	Gly	Gln	Gln	Gly 265	Arg	Gly	Arg	Ala	Gly 270
Ile	Glu	Ala	Ser	Leu 275	Asp	Val	Gln	Tyr	Leu 280	Met	Ser	Ala	Gly	Ala 285
Asn	Ile	Ser	Thr	Trp 290	Val	Tyr	Ser	Ser	Pro 295	Gly	Arg	His	Glu	Gly 300
Gln	Glu	Pro	Phe	Leu 305	Gln	Trp	Leu	Met	Leu 310	Leu	Ser	Asn	Glu	Ser 315
Ala	Leu	Pro	His	Val 320	His	Thr	Val	Ser	Tyr 325	Gly	Asp	Asp	Glu	Asp 330
Ser	Leu	Ser	Ser	Ala 335	Tyr	Ile	Gln	Arg	Val 340	Asn	Thr	Glu	Leu	Met 345
Lys	Ala	Ala	Ala	Arg 350	Gly	Leu	Thr	Leu	Leu 355	Phe	Ala	Ser	Gly	Asp 360
Ser	Gly	Ala	Gly	Cys 365	Trp	Ser	Val	Ser	Gly 370	Arg	His	Gln	Phe	Arg 375
Pro	Thr	Phe	Pro	Ala 380	Ser	Ser	Pro	Tyr	Val 385	Thr	Thr	Val	Gly	Gly 390
Thr	Ser	Phe	Gln	Glu 395	Pro	Phe	Leu	Ile	Thr 400	Asn	Glu	Ile	Val	Asp 405
Tyr	Ile	Ser	Gly	Gly 410	Gly	Phe	Ser	Asn	Val 415	Phe	Pro	Arg	Pro	Ser 420
Tyr	Gln	Glu	Glu	Ala 425	Val	Thr	Lys	Phe	Leu 430	Ser	Ser	Ser	Pro	His 435
Leu	Pro	Pro	Ser	Ser 440	Tyr	Phe	Asn	Ala	Ser 445	Gly	Arg	Ala	Tyr	Pro 450
Asp	Val	Ala	Ala	Leu 455	Ser	Asp	Gly	Tyr	Trp 460	Val	Val	Ser	Asn	Arg 465
Val	Pro	Ile	Pro	Trp 470	Val	Ser	Gly	Thr	Ser 475	Ala	Ser	Thr	Pro	Val 480
Phe	Gly	Gly	Ile	Leu 485	Ser	Leu	Ile	Asn	Glu 490	His	Arg	Ile	Leu	Ser 495
Gly	Arg	Pro	Pro	Leu 500	Gly	Phe	Leu	Asn	Pro 505	Arg	Leu	Tyr	Gln	Gln 510
His	Gly	Ala	Gly	Leu 515	Phe	Asp	Val	Thr	Arg 520	Gly	Cys	His	Glu	Ser 525
Cys	Leu	Asp	Glu	Glu 530	Val	Glu	Gly	Gln	Gly 535	Phe	Cys	Ser	Gly	Pro 540

Cys

<210> 260

<211> 1638

<212> DNA

<213> Homo Sapien

<400> 260

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ctattggatt aaaggaaact acctggattg tagggagggg tgacacagtg 1250
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caaactttga ttttatttc atctgaactt gtttcaaaga tttatattaa 1600
atatttggca tacaagagat atgaaaaaaa aaaaaaaa 1638

<210> 261

<211> 383

<212> PRT

<213> Homo Sapien

<400> 261

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Thr Trp Pro Ala Tyr Arg Leu Pro Val Val Leu Pro Gln Ser Thr
35 40 45

Leu Asn Leu Ala Lys Pro Asp Phe Gly Ala Glu Ala Lys Leu Glu 50 55 60

Val Ser Ser Ser Cys Gly Pro Gln Cys His Lys Gly Thr Pro Leu 65 70 75

Pro Thr Tyr Glu Glu Ala Lys Gln Tyr Leu Ser Tyr Glu Thr Leu 80 85 90

Tyr Ala Asn Gly Ser Arg Thr Glu Thr Gln Val Gly Ile Tyr Ile \$95\$ 100 105

Leu Ser Ser Ser Gly Asp Gly Ala Gln His Arg Asp Ser Gly Ser 110 115 120

Ser Gly Lys Ser Arg Arg Lys Arg Gln Ile Tyr Gly Tyr Asp Ser 125 130 135

Arg Phe Ser Ile Phe Gly Lys Asp Phe Leu Leu Asn Tyr Pro Phe 140 145 150

Ser Thr Ser Val Lys Leu Ser Thr Gly Cys Thr Gly Thr Leu Val

Ala Glu Lys His Val Leu Thr Ala Ala His Cys Ile His Asp Gly
170 175 180

Lys Thr Tyr Val Lys Gly Thr Gln Lys Leu Arg Val Gly Phe Leu 1.85 Lys Pro Lys Phe Lys Asp Gly Gly Arg Gly Ala Asn Asp Ser Thr 200 Ser Ala Met Pro Glu Gln Met Lys Phe Gln Trp Ile Arg Val Lys 220 215 Arg Thr His Val Pro Lys Gly Trp Ile Lys Gly Asn Ala Asn Asp Ile Gly Met Asp Tyr Asp Tyr Ala Leu Leu Glu Leu Lys Lys Pro 245 His Lys Arg Lys Phe Met Lys Ile Gly Val Ser Pro Pro Ala Lys 260 270 Gln Leu Pro Gly Gly Arg Ile His Phe Ser Gly Tyr Asp Asn Asp 275 Arg Pro Gly Asn Leu Val Tyr Arg Phe Cys Asp Val Lys Asp Glu 290 Thr Tyr Asp Leu Leu Tyr Gln Gln Cys Asp Ala Gln Pro Gly Ala 305 Ser Gly Ser Gly Val Tyr Val Arg Met Trp Lys Arg Gln Gln Gln 320 330 Lys Trp Glu Arg Lys Ile Ile Gly Ile Phe Ser Gly His Gln Trp 335 340 Val Asp Met Asn Gly Ser Pro Gln Asp Phe Asn Val Ala Val Arg 350 355 360 Ile Thr Pro Leu Lys Tyr Ala Gln Ile Cys Tyr Trp Ile Lys Gly 365 Asn Tyr Leu Asp Cys Arg Glu Gly 380

<210> 262

<211> 1378

<212> DNA

<213> Homo Sapien

<400> 262

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ctgggtgatc actgctgccc actgtttcaa ggacaacctg aacaaccat 350 acctgttctc tgtgctgctg ggggcctggc agctggggaa ccctggctct 400 cggtcccaga aggtgggtgt tgcctgggtg gagccccacc ctgtgtattc 450 ctggaaggaa ggtgcctgtg cagacattgc cctggtgcgt ctcgagcgct 500 ccatacagtt ctcagagegg gtcctgccca tctgcctacc tgatgcctct 550 atccacctcc ctccaaacac ccactgctgg atctcaggct gggggagcat 600 ccaagatgga gttcccttgc cccaccctca gaccctgcag aagctgaagg 650 tteetateat egaeteggaa gtetgeagee atetgtaetg geggggagea 700 ggacagggac ccatcactga ggacatgctg tgtgccggct acttggaggg 750 ggagcgggat gcttgtctgg gcgactccgg gggcccctc atgtgccagg 800 tggacggcgc ctggctgctg gccggcatca tcagctgggg cgagggctgt 850 gccgagcgca acaggcccgg ggtctacatc agcctctctg cgcaccgctc 900 ctgggtggag aagatcgtgc aaqgggtgca gctccgcqqg cgcqctcaqg 950 ggggtgggc cctcagggca ccgagccagg gctctggggc cgccgcgcgc 1000 tectagggeg cagegggaeg eggggetegg atetgaaagg eggeeagate 1050 cacatetgga tetggatetg eggeggeete gggeggttte eeeegeegta 1100 aataggetea tetaceteta eetetggggg eeeggaegge tgetgeggaa 1150 aggaaaccc ctccccgacc cgcccgacgg cctcaggccc ccctccaagg 1200 catcaggccc cgcccaacgg cctcatgtcc ccgccccac gacttccggc 1250 cccgcccccg ggccccagcg cttttgtgta tataaatgtt aatgattttt 1300 ataggtattt gtaaccctgc ccacatatct tatttattcc tccaatttca 1350 ataaattatt tattctccaa aaaaaaaa 1378

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- <211> 317
- <212> PRT
- <213> Homo Sapien
- <400> 263
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- Gly Thr Phe Thr Ser Leu Leu Leu Leu Ala Ser Thr Ala Ile Leu 20 25 30
- Asn Ala Ala Arg Ile Pro Val Pro Pro Ala Cys Gly Lys Pro Gln
 35 40 45

Gln	Leu	Asn	Arg	Val 50	Val	Gly	Gly	Glu	Asp 55	Ser	Thr	Asp	Ser	Glu 60
Trp	Pro	Trp	Ile	Val 65	Ser	Ile	Gln	Lys	Asn 70	Gly	Thr	His	His	Cys 75
Ala	Gly	Ser	Leu	Leu 80	Thr	Ser	Arg	Trp	Val 85	Ile	Thr	Ala	Ala	His 90
Cys	Phe	Lys	Asp	Asn 95	Leu	Asn	Lys	Pro	Tyr 100	Leu	Phe	Ser	Val	Leu 105
Leu	Gly	Ala	Trp	Gln 110	Leu	Gly	Asn	Pro	Gly 115	Ser	Arg	Ser	Gln	Lys 120
Val	Gly	Val	Ala	Trp 125	Val	Glu	Pro	His	Pro 130	Val	Tyr	Ser	Trp	Lys 135
Glu	Gly	Ala	Cys	Ala 140	Asp	Ile	Ala	Leu	Val 145	Arg	Leu	Glu	Arg	Ser 150
Ile	Gln	Phe	Ser	Glu 155	Arg	Val	Leu	Pro	Ile 160	Cys	Leu	Pro	Asp	Ala 165
Ser	Ile	His	Leu	Pro 170	Pro	Asn	Thr	His	Cys 175	Trp	Ile	Ser	Gly	Trp 180
Gly	Ser	Ile	Gln	Asp 185	Gly	Val	Pro	Leu	Pro 190	His	Pro	Gln	Thr	Leu 195
Gln	Lys	Leu	Lys	Val 200	Pro	Ile	Ile	Asp	Ser 205	Glu	Val	Cys	Ser	His 210
Leu	Tyr	Trp	Arg	Gly 215	Ala	Gly	Gln	Gly	Pro 220	Ile	Thr	Glu	Asp	Met 225
Leu	Cys	Ala	Gly	Tyr 230	Leu	Glu	Gly	Glu	Arg 235	Asp	Ala	Cys	Leu	Gly 240
	Ser			245					250				_	255
Leu	Ala	Gly	Ile	Ile 260	Ser	Trp	Gly	Glu	Gly 265	Cys	Ala	Glu	Arg	Asn 270
Arg	Pro	Gly	Val	Tyr 275	Ile	Ser	Leu	Ser	Ala 280	His	Arg	Ser	Trp	Val 285
Glu	Lys	Ile	Val	Gln 290	Gly	Val	Gln	Leu	Arg 295	Gly	Arg	Ala	Gln	Gly 300
Gly	Gly	Ala	Leu	Arg 305	Ala	Pro	Ser	Gln	Gly 310	Ser	Gly	Ala	Ala	Ala 315

Arg Ser

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<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 264
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<210> 265
<211> 19
<212> DNA
<213> Artificial Sequence
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<400> 265
 gcagaggtgt ctaaggttg 19
<210> 266
<211> 24
<212> DNA
<213> Artificial Sequence
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<223> Synthetic Oligonucleotide Probe
<400> 266
 agctctagac caatgccagc ttcc 24
<210> 267
<211> 45
<212> DNA
<213> Artificial Sequence
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<400> 267
 gccaccaact cctgcaagaa cttctcagaa ctgcccctgg tcatg 45
<210> 268
<211> 25
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<400> 268
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<400> 269
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<223> Synthetic Oligonucleotide Probe
<400> 270
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<210> 271
<211> 26
<212> DNA
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<400> 271
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<400> 272
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<212> DNA
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<223> Synthetic Oligonucleotide Probe
<400> 273
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<211> 24
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<400> 274

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<211> 21
<212> DNA
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<223> Synthetic Oligonucleotide Probe
<400> 276
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<210> 277
<211> 18
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<400> 277
ggctatgaca gcaggttc 18
<210> 278
<211> 18
<212> DNA
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<223> Synthetic Oligonucleotide Probe
<400> 278
 tgacaatgac cgaccagg 18
<210> 279
<211> 24
<212> DNA
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<223> Synthetic Oligonucleotide Probe
<400> 279
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<210> 280
<211> 45
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<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 280
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<210> 281
<211> 34
<212> DNA
<213> Artificial Sequence
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<223> Synthetic Oligonucleotide Probe
<400> 281
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<210> 282
<211> 61
<212> DNA
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<220>
<223> Synthetic Oligonucleotide Probe
<400> 282
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tgccaggtgg a 61
<210> 283
<211> 119
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 283
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atgctgtgtg ccggctact 119
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<211> 1875
<212> DNA
<213> Homo Sapien
<400> 284
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- <212> PRT
- <213> Homo Sapien
- <400> 292
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 Leu Ala Leu Gln Leu Leu Val Val Ala Gly Leu Val Arg Ala Gln
 35 40 45

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Thr	Asn	Thr	Arg	Leu 80	Leu	Asn	Leu	His	Glu 85	Asn	Gln	Ile	Gln	Ile 90
Ile	Lys	Val	Asn	Ser 95	Phe	Lys	His	Leu	Arg 100	His	Leu	Glu	Ile	Leu 105
Gln	Leu	Ser	Arg	Asn 110	His	Ile	Arg	Thr	Ile 115	Glu	Ile	Gly	Ala	Phe 120
Asn	Gly	Leu	Ala	Asn 125	Leu	Asn	Thr	Leu	Glu 130	Leu	Phe	Asp	Asn	Arg 135
Leu	Thr	Thr	Ile	Pro 140	Asn	Gly	Ala	Phe	Val 145	Tyr	Leu	Ser	Lys	Leu 150
Lys	Glu	Leu	Trp	Leu 155	Arg	Asn	Asn	Pro	Ile 160	Glu	Ser	Ile	Pro	Ser 165
Tyr	Ala	Phe	Asn	Arg 170	Ile	Pro	Ser	Leu	Arg 175	Arg	Leu	Asp	Leu	Gly 180
Glu	Leu	Lys	Arg	Leu 185	Ser	Tyr	Ile	Ser	Glu 190	Gly	Ala	Phe	Glu	Gly 195
Leu	Ser	Asn	Leu	Arg 200	Tyr	Leu	Asn	Leu	Ala 205	Met	Cys	Asn	Leu	Arg 210
Glu	Ile	Pro	Asn	Leu 215	Thr	Pro	Leu	Ile	Lys 220	Leu	Asp	Glu	Leu	Asp 225
				215		Pro Ser			220					225
Leu	Ser.	Gly	Asn	215 His 230	Leu		Ala	Ile	220 Arg 235	Pro	Gly	Ser	Phe	225 Gln 240
Leu	Ser.	Gly Met	Asn His	215 His 230 Leu 245	Leu Gln	Ser	Ala Leu	Ile Trp	220 Arg 235 Met 250	Pro Ile	Gly Gln	Ser Ser	Phe Gln	225 Gln 240 Ile 255
Leu	Ser Leu Val	Gly Met Ile	Asn His Glu	215 His 230 Leu 245 Arg 260	Leu Gln Asn	Ser Lys	Ala Leu Phe	Ile Trp Asp	220 Arg 235 Met 250 Asn 265	Pro Ile Leu	Gly Gln Gln	Ser Ser	Phe Gln Leu	225 Gln 240 Ile 255 Val 270
Leu Gly Gln	Ser Leu Val	Gly Met Ile Asn	Asn His Glu Leu	215 His 230 Leu 245 Arg 260 Ala 275	Leu Gln Asn His	Ser Lys Ala	Ala Leu Phe Asn	Ile Trp Asp Leu	220 Arg 235 Met 250 Asn 265 Thr 280	Pro Ile Leu Leu	Gly Gln Gln Leu	Ser Ser Ser	Phe Gln Leu His	225 Gln 240 Ile 255 Val 270 Asp 285
Leu Gly Gln Glu Leu	Ser Leu Val Ile	Gly Met Ile Asn Thr	Asn His Glu Leu Pro	215 His 230 Leu 245 Arg 260 Ala 275 Leu 290	Leu Gln Asn His	Ser Lys Ala Asn	Ala Leu Phe Asn Leu	Ile Trp Asp Leu Glu	220 Arg 235 Met 250 Asn 265 Thr 280 Arg 295	Pro Ile Leu Leu Ile	Gly Gln Gln Leu	Ser Ser Pro	Phe Gln Leu His	225 Gln 240 Ile 255 Val 270 Asp 285 His 300
Leu Gly Gln Glu Leu Asn	Ser Leu Val Ile Phe	Gly Met Ile Asn Thr	Asn His Glu Leu Pro Asn	215 His 230 Leu 245 Arg 260 Ala 275 Leu 290 Cys 305	Leu Gln Asn His Asn	Ser Lys Ala Asn His	Ala Leu Phe Asn Leu	Ile Trp Asp Leu Glu Ile	220 Arg 235 Met 250 Asn 265 Thr 280 Arg 295 Leu 310	Pro Ile Leu Leu Ile	Gly Gln Gln Leu His	Ser Ser Pro Leu Ser	Phe Gln Leu His Trp	225 Gln 240 Ile 255 Val 270 Asp 285 His 300 Trp 315

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Ala	Asp	Leu	Asn	Val 365	Thr	Glu	Gly	Met	Ala 370	Ala	Glu	Leu	Lys	Cys 375
Arg	Ala	Ser	Thr	Ser 380	Leu	Thr	Ser	Val	Ser 385	Trp	Ile	Thr	Pro	Asn 390
Gly	Thr	Val	Met	Thr 395	His	Gly	Ala	Tyr	Lys 400	Val	Arg	Ile	Ala	Val 405
Leu	Ser	Asp	Gly	Thr 410	Leu	Asn	Phe	Thr	Asn 415	Val	Thr	Val	Gln	Asp 420
Thr	Gly	Met	Tyr	Thr 425	Cys	Met	Val	Ser	Asn 430	Ser	Val	Gly	Asn	Thr 435
Thr	Ala	Ser	Ala	Thr 440	Leu	Asn	Val	Thr	Ala 445	Ala	Thr	Thr	Thr	Pro 450
Phe	Ser	Tyr	Phe	Ser 455	Thr	Val	Thr	Val	Glu 460	Thr	Met	Glu	Pro	Ser 465
Gln	Asp	Glu	Ala	Arg 470	Thr	Thr	Asp	Asn	Asn 475	Val	Gly	Pro	Thr	Pro 480
Val	Val	Asp	Trp	Glu 485	Thr	Thr	Asn	Val	Thr 490	Thr	Ser	Leu	Thr	Pro 495
Gln	Ser	Thr	Arg	Ser 500	Thr	Glu	Lys	Thr	Phe 505	Thr	Ile	Pro	Val	Thr 510
Asp	Ile	Asn	Ser	Gly 515	Ile	Pro	Gly	Ile	Asp 520	Glu	Val	Met	Lys	Thr 525
Thr	Lys	Ile	Ile	Ile 530	Gly	Cys	Phe	Val	Ala 535	Ile	Thr	Leu	Met	Ala 540
Ala	Val	Met	Leu	Val 545	Ile	Phe	Tyr	Lys	Met 550	Arg	Lys	Gln	His	His 555
Arg	Gln	Asn	His	His 560	Ala	Pro	Thr	Arg	Thr 565	Val	Glu	Ile	Ile	Asn 570
Val	Asp	Asp	Glu	Ile 575	Thr	Gly	Asp	Thr	Pro 580	Met	Glu	Ser	His	Leu 585
Pro	Met	Pro	Ala	Ile 590	Glu	His	Glu	His	Leu 595	Asn	His	Tyr	Asn	Ser 600
Tyr	Lys	Ser	Pro	Phe 605	Asn	His	Thr	Thr	Thr 610	Val	Asn	Thr	Ile	Asn 615
Ser	Ile	His	Ser	Ser 620	Val	His	Glu	Pro	Leu 625	Leu	Ile	Arg	Met	Asn 630
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- <210> 293
- <211> 4053
- <212> DNA
- <213> Homo Sapien
- <400> 293

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<212> PRT

<213> Homo Sapien

<400> 294

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Gly	Arg	Gly	Glu	Leu 35	Gly	Gln	Pro	Ser	Gly 40	Val	Ala	Ala	Glu	Arg 45
Pro	Cys	Pro	Thr	Thr 50	Cys	Arg	Cys	Leu	Gly 55	Asp	Leu	Leu	Asp	Cys 60
Ser	Arg	Lys	Arg	Leu 65	Ala	Arg	Leu	Pro	Glu 70	Pro	Leu	Pro	Ser	Trp 75
Val	Ala	Arg	Leu	Asp 80	Leu	Ser	His	Asn	Arg 85	Leu	Ser	Phe	Ile	Lys 90
Ala	Ser	Ser	Met		His	Leu	Gln	Ser		Arg	Glu	Val	Lys	
Asn	Asn	Asn	Glu	Leu 110	Glu	Thr	Ile	Pro	Asn 115	Leu	Gly	Pro	Val	Ser 120
Ala	Asn	Ile	Thr	Leu 125	Leu	Ser	Leu	Ala	Gly 130	Asn	Arg	Ile	Val	Glu 135
Ĭle	Leu	Pro	Glu	His 140	Leu	Lys	Glu	Phe	Gln 145	Ser	Leu	Glu	Thr	Leu 150
Asp	Leu	Ser	Ser	Asn 155	Asn	Ile	Ser	Glu	Leu 160	Gln	Thr	Ala	Phe	Pro 165
Ala	Leu	Gln	Leu	Lys 170	Tyr	Leu	Tyr	Leu	Asn 175	Ser	Asn	Arg	Val	Thr 180
Ser	Met	Glu	Pro	Gly 185	Tyr	Phe	Asp	Asn	Leu 190	Ala	Asn	Thr	Leu	Leu 195
Val	Leu	Lys	Leu	Asn 200	Arg	Asn	Arg	Ile	Ser 205	Ala	Ile	Pro	Pro	Lys 210
Met	Phe	Lys	Leu	Pro 215	Gln	Leu	Gln	His	Leu 220	Glu	Leu	Asn	Arg	Asn 225
Lys	Ile	Lys	Asn	Val 230	Asp	Gly	Leu	Thr	Phe 235	Gln	Gly	Leu	Gly	Ala 240
Leu	Lys	Ser	Leu	Lys 245	Met	Gln	Arg	Asn	Gly 250	Val	Thr	Lys	Leu	Met 255
Asp	Gly	Ala	Phe	Trp 260	Gly	Leu	Ser	Asn	Met 265	Glu	Ile	Leu	Gln	Leu 270
Asp	His	Asn	Asn	Leu 275	Thr	Glu	Ile	Thr	Lys 280	Gly	Trp	Leu	Tyr	Gly 285
Leu	Leu	Met	Leu	Gln 290	Glu	Leu	His	Leu	Ser 295	Gln	Asn	Ala	Ile	Asn 300

Arg	Ile	Ser	Pro	Asp 305	Ala	Trp	Glu	Phe	Cys 310	Gln	Lys	Leu	Ser	Glu 315
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Phe	Leu	Gly	Leu	Ser 335	Leu	Leu	Asn	Thr	Leu 340	His	Ile	Gly	Asn	Asn 345
Arg	Val	Ser	Tyr	Ile 350	Ala	Asp	Cys	Ala	Phe 355	Arg	Gly	Leu	Ser	Ser 360
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Glu	Asp	Met	Asn	Gly 380	Ala	Phe	Ser	Gly	Leu 385	Asp	Lys	Leu	Arg	Arg 390
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Phe	Thr	Gly	Leu	Asp 410	Ala	Leu	Glu	His	Leu 415	Asp	Leu	Ser	Asp	Asn 420
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Phe	Pro	Lys	Pro	Gln 500	Ile	Thr	Val	Gln	Pro 505	Glu	Thr	Gln	Ser	Ala 510
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Gly	Gly	Glu	Val	Met 560	Glu	Tyr	Thr	Thr	Ile 565	Leu	Arg	Leu	Arg	Glu 570
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Phe	Phe	Ile	Val	Asp 665	Val	Lys	Ile	Glu	Asp 670	Ile	Gly	Val	Tyr	Ser 675
Cys	Thr	Ala	Gln	Asn 680	Ser	Ala	Gly	Ser	Ile 685	Ser	Ala	Asn	Ala	Thr 690
Leu	Thr	Val	Leu	Glu 695	Thr	Pro	Ser	Phe	Leu 700	Arg	Pro	Leu	Leu	Asp 705
Arg	Thr	Val	Thr	Lys 710	Gly	Glu	Thr	Ala	Val 715	Leu	Gln	Cys	Ile	Ala 720
Gly	Gly	Ser	Pro	Pro 725	Pro	Lys	Leu	Asn	Trp 730	Thr	Lys	Asp	Asp	Ser 735
Pro	Leu	Val	Val	Thr 740	Glu	Arg	His	Phe	Phe 745	Ala	Ala	Gly	Asn	Gln 750
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Thr	Cys	Glu	Met	Ser 770	Asn	Thr	Leu	Gly	Thr 775	Glu	Arg	Gly	Asn	Val 780
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Tyr	Val	Ser	Ser	Glu 875	Ser	Gly	Ser	His	His 880	Gln	Phe	Val	Thr	Ser 885
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Glu	Gly	Pro	Gly	Met L010	Lys	Asn	Leu		Leu 1015	Asn	Lys	Ser		Leu 1020
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Ser	Phe	Met	Gly 1	Thr L040	Phe	Gly	Lys		Leu 1045	Arg	Arg	Pro		Leu 1050
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Ala	Phe	Tyr	Leu 1	Lys 1070	Ala	His	Ser		Pro .075	Asp	Leu	Asp		Gly 1080
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Gln Pro Arg Cys Lys His Gly Glu Cys Ile Gly Pro Asn Lys Cys
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Lys Cys His Pro Gly Tyr Ala Gly Lys Thr Cys Asn Gln Asp Leu 80 85 90

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- Gly Val Arg Gly Gln Ala Leu Tyr Leu Pro Val His Tyr Gly Phe
 35 40 45
- His Thr Pro Ala Ser Asp Ile Gln Ile Ile Trp Leu Phe Glu Arg
 50 55 60
- Pro His Thr Met Pro Lys Tyr Leu Leu Gly Ser Val Asn Lys Ser
 65 70 75
- Val Val Pro Asp Leu Glu Tyr Gln His Lys Phe Thr Met Met Pro 80 85 90
- Pro Asn Ala Ser Leu Leu Ile Asn Pro Leu Gln Phe Pro Asp Glu
 95 100 105

Gly	Asn	Tyr	Ile	Val 110	Lys	Val	Asn	Ile	Gln 115	Gly	Asn	Gly	Thr	Leu 120
Ser	Ala	Ser	Gln	Lys 125	Ile	Gln	Val	Thr	Val 130	Asp	Asp	Pro	Val	Thr 135
Lys	Pro	Val	Val	Gln 140	Ile	His	Pro	Pro	Ser 145	Gly	Ala	Val	Glu	Tyr 150
Val	Gly	Asn	Met	Thr 155	Leu	Thr	Cys	His	Val 160	Glu	Gly	Gly	Thr	Arg 165
Leu	Ala	Tyr	Gln	Trp 170	Leu	Lys	Asn	Gly	Arg 175	Pro	Val	His	Thr	Ser 180
Ser	Thr	Tyr	Ser	Phe 185	Ser	Pro	Gln	Asn	Asn 190	Thr	Leu	His	Ile	Ala 195
Pro	Val	Thr	Lys	Glu 200	Asp	Ile	Gly	Asn	Tyr 205	Ser	Cys	Leu	Val	Arg 210
Asn	Pro	Val	Ser	Glu 215	Met	Glu	Ser	Asp	Ile 220	Ile	Met	Pro	Ile	Ile 225
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Lys	Val	Gly	Glu	Val 245	Phe	Thr	Val	Asp	Leu 250	Gly	Glu	Ala	Ile	Leu 255
Phe	Asp	Cys	Ser	Ala 260	Asp	Ser	His	Pro	Pro 265	Asn	Thr	Tyr	Ser	Trp 270
Ile	Arg	Arg	Thr	Asp 275	Asn	Thr	Thr	Tyr	Ile 280	Ile	Lys	His	Gly	Pro 285
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Ala	Gln	Lys	Gly	Lys 335	Ser	Leu	Ser	Pro	Leu 340	Ala	Ser	Ile	Thr	Gly 345
Ile	Ser	Leu	Phe	Leu 350	Ile	Ile	Ser,	Met	Cys 355	Leu	Leu	Phe	Leu	Trp 360
Lys	Lys	Tyr	Gln	Pro 365	Tyr	Lys	Val	Ile	Lys 370	Gln	Lys	Leu	Glu	Gly 375
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tggatgatag aattttatgc cccgtggtgc cctgcttgtc aaaatcttca 200
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accggaatgg gaaagttttg ctgaatgggg agaagatctt gaggttaata 250

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Val Arg Val Ile Thr Asp Glu Asn Trp Arg Glu Leu Leu Glu Gly
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Asp Trp Met Ile Glu Phe Tyr Ala Pro Trp Cys Pro Ala Cys Gln
50 55 60

Asn Leu Gln Pro Glu Trp Glu Ser Phe Ala Glu Trp Gly Glu Asp
65 70 75

Leu Glu Val Asn Ile Ala Lys Val Asp Val Thr Glu Gln Pro Gly
80 85 90

Leu Ser Gly Arg Phe Ile Ile Thr Ala Leu Pro Thr Ile Tyr His
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Cys Lys Asp Gly Glu Phe Arg Arg Tyr Gln Gly Pro Arg Thr Lys 110 115 120

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<212> PRT

<213> Homo Sapien

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Thr Asp Leu Ile Tyr Ala Glu Lys Glu Leu Val Gln Ser Leu Lys
35 40 45

Glu Tyr Ile Leu Val Glu Glu Ala Lys Leu Ser Lys Ile Lys Ser
50 55 60

Trp Ala Asn Lys Met Glu Ala Leu Thr Ser Lys Ser Ala Ala Asp
65 70 75

Ala Glu Gly Tyr Leu Ala His Pro Val Asn Ala Tyr Lys Leu Val 80 85 90

Lys Arg Leu Asn Thr Asp Trp Pro Ala Leu Glu Asp Leu Val Leu 95 100 105

Gln Asp Ser Ala Ala Gly Phe Ile Ala Asn Leu Ser Val Gln Arg

Gln Phe Phe Pro Thr Asp Glu Asp Glu Ile Gly Ala Ala Lys Ala 125 130 135

Leu Met Arg Leu Gln Asp Thr Tyr Arg Leu Asp Pro Gly Thr Ile 140 145 150

Ser Arg Gly Glu Leu Pro Gly Thr Lys Tyr Gln Ala Met Leu Ser 155 160 165

Val Asp Asp Cys Phe Gly Met Gly Arg Ser Ala Tyr Asn Glu Gly
170 175 180

Asp Tyr Tyr His Thr Val Leu Trp Met Glu Gln Val Leu Lys Gln
185 190 195

Leu Asp Ala Gly Glu Glu Ala Thr Thr Lys Ser Gln Val Leu 200 205 210

Asp Tyr Leu Ser Tyr Ala Val Phe Gln Leu Gly Asp Leu His Arg 215 220 225

Ala Leu Glu Leu Thr Arg Arg Leu Leu Ser Leu Asp Pro Ser His
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Glu Arg Ala Gly Gly Asn Leu Arg Tyr Phe Glu Gln Leu Leu Glu

Glu Glu Arg Glu Lys Thr Leu Thr Asn Gln Thr Glu Ala Glu Leu

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	Arg	Pro	Phe	Asp	Ser 440	Gly	Leu	Lys	Thr	Glu 445	Gly	Asn	Arg	Leu	Ala 450
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	Val	Phe	Pro	Asp	Leu 470	Gly	Ala	Ala	Ile	Trp 475	Pro	Lys	Lys	Gly	Thr 480
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	Arg	Thr	Arg	His	Ala 500	Ala	Cys	Pro	Val	Leu 505	Val	Gly	Cys	Lys	Trp 510
	Val	Ser	Asn	Lys	Trp 515	Phe	His	Glu	Arg	Gly 520	Gln	Glu	Phe	Leu	Arg 525
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<211> 772

<212> PRT

<213> Homo Sapien

<400> 339

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Tyr	Arg	Asp	Pro	Asn 80	Lys	Pro	Tyr	Lys	Lys 85	Val	Leu	Arg	Thr	Arg 90
Tyr	Ile	Gln	Thr	Glu 95	Leu	Gly	Ser	Arg	Glu 100	Arg	Leu	Leu	Val	Ala 105
Val	Leu	Thr	Ser	Arg 110	Ala	Thr	Leu	Ser	Thr 115	Leu	Ala	Val	Ala	Val 120
Asn	Arg	Thr	Val	Ala 125	His	His	Phe	Pro	Arg 130		Leu	Tyr	Phe	Thr 135
Gly	Gln	Arg	Gly	Àla 140	Arg	Ala	Pro	Ala	Gly 145	Met	Gln	Val	Val	Ser 150
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Pro	Ser	Gln	Val	Arg 575	Leu	Met	Asp	Val	Val 580	Ser	Lys	Lys	His.	Pro 585
Val	Asp	Thr	Leu	Phe 590	Phe	Leu	Thr	Thr	Val 595	Trp	Thr	Arg	Pro	Gly 600
Pro	Glu	Val	Leu	Asn	Arg	Cys	Arg	Met	Asn	Ala	Ile	Ser	Gly	Trp

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Met	Glu	Leu	Ser	Lys 65	Ser	Phe	Arg	Val	Tyr 70	Cys	Ile	Ile	Leu	Val 75
Lys	Pro	Lys	Asp	Val 80	Ser	Leu	Trp	Ala	Ala 85	Val	Lys	Glu	Thr	Trp 90
Thr	Lys	His	Cys	Asp 95	Lys	Ala	Glu	Phe	Phe 100	Ser	Ser	Glu	Asn	Val 105
Lys	Val	Phe	Glu	Ser 110	Ile	Asn	Met	Asp	Thr 115	Asn	Asp	Met	Trp	Leu 120
Met	Met	Arg	Lys	Ala 125	Tyr	Lys	Tyr	Ala	Phe 130	Asp	Lys	Tyr	Arg	Asp 135
Gln	Tyr	Asn	Trp	Phe 140	Phe	Leu	Ala	Arg	Pro 145	Thr	Thr	Phe	Ala	Ile 150
Ile	Glu	Asn	Leu	Lys 155	Tyr	Phe	Leu	Leu	Lys 160	Lys	Asp	Pro	Ser	Gln 165
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Gly	Gly	Met	Ile	Trp 215	Lys	Ile	Ser	Glu	Asp 220	Lys	Gln	Leu	Ala	Val 225
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Cys	Ser	Asp	Met	Ala 275	Val	Thr	Phe	Asn	Gly 280	Leu	Thr	Pro	Asn	Gln 285
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gagagactet teaaaceeaa ggeaaaattg gaatgatgta acetgtttee 700
tcaattattt tcggatttgt gaaatggtag gaataaatcc tttgaacaaa 750
ggaaaatctc tttaagaaca gaaggcacaa ctcaaatgtg taaagaagga 800
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agagcaagaa catggccaca cccaccgccc cacacgagaa atttgtgcgc 850

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- <210> 377
- <211> 219
- <212> PRT
- <213> Homo Sapien

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- Cys Phe Ser Ser Gln Met Phe Leu Trp Thr Val Ala Gly Ile Pro 20 25 30
- Ile Leu Phe Leu Ser Ala Cys Phe Ile Thr Arg Cys Val Val Thr
 35 40 45
- Phe Arg Ile Phe Gln Thr Cys Asp Glu Lys Lys Phe Gln Leu Pro
 50 55 60
- Glu Asn Phe Thr Glu Leu Ser Cys Tyr Asn Tyr Gly Ser Gly Ser
 65 70 75
- Val Lys Asn Cys Cys Pro Leu Asn Trp Glu Tyr Phe Gln Ser Ser 80 85 90
- Cys Tyr Phe Phe Ser Thr Asp Thr Ile Ser Trp Ala Leu Ser Leu 95 100 105
- Lys Asn Cys Ser Ala Met Gly Ala His Leu Val Val Ile Asn Ser 110 115 120
- Gln Glu Glu Glu Phe Leu Ser Tyr Lys Lys Pro Lys Met Arg 125 130 135
- Glu Phe Phe Ile Gly Leu Ser Asp Gln Val Val Glu Gly Gln Trp 140 145 150
- Gln Trp Val Asp Gly Thr Pro Leu Thr Lys Ser Leu Ser Phe Trp 155 160 165
- Asp Val Gly Glu Pro Asn Asn Ile Ala Thr Leu Glu Asp Cys Ala 170 175 180
- Thr Met Arg Asp Ser Ser Asn Pro Arg Gln Asn Trp Asn Asp Val 185 190 195
- Thr Cys Phe Leu Asn Tyr Phe Arg Ile Cys Glu Met Val Gly Ile 200 205 210
- Asn Pro Leu Asn Lys Gly Lys Ser Leu 215
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catcaagcgc ctctacca 18
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gccgctgtcc actgcag 17
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<210> 412
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<210> 415
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Ala Val Asn Leu Lys Ser Ser Asn Arg Thr Pro Val Val Gln Glu
35 40 45

Phe Glu Ser Val Glu Leu Ser Cys Ile Ile Thr Asp Ser Gln Thr
50 55 60

Ser Asp Pro Arg Ile Glu Trp Lys Lys Ile Gln Asp Glu Gln Thr
65 70 75

Thr Tyr Val Phe Phe Asp Asn Lys Ile Gln Gly Asp Leu Ala Gly 80 85 90

Arg Ala Glu Ile Leu Gly Lys Thr Ser Leu Lys Ile Trp Asn Val

<210> 423

<211> 310

<212> PRT

<213> Homo Sapien

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Gln	Val	Lys	Pro	Val 140	Thr	Pro	Val	Cys	Arg 145	۷a - ُ.	Pro	Lys	Ala	Val 150
Pro	Val	Gly	Lys	Met 155	Ala	Thr	Leu	His	Cys 160	Gln	Glu	Ser	Glu	Gly 165
His	Pro	Arg	Pro	His 170	Tyr	Ser	Trp	Tyr	Arg 175	Asn	Asp	Val	Pro	Leu 180
Pro	Thr	Asp	Ser	Arg 185	Ala	Asn	Pro	Arg	Phe 190	Arg	Asn	Ser	Ser	Phe 195
His	Leu	Asn	Ser	Glu 200	Thr	Gly	Thr	Leu	Val 205	Phe	Thr	Ala	Val	His 210
Lys	Asp	Asp	Ser	Gly 215	Gln	Tyr	Tyr	Cys	11e 220	Ala	Ser	Asn	Asp	Ala 225
Gly	ser Ser	Ala	Arg	Cys 230	Glu	Glu	Gln	Glu	Met 235	Glu	Val	Tyr	Asp	Leu 240
Asr	ılle	Gly	gly	7 Ile 245	Ile	gly	Gly	Val	Leu 250	Val	Val	Leu	Ala	Val 255
Let	ı Ala	Lev	ı Ile	260	Leu	ı Gly	, Ile	c Cys	265	Ala	Tyr	Arg	Arg	Gly 270
Ту	r Phe	e Ile	e Asr	n Asr 275	ı Lys	s Glr	a Asp	Gly	7 Glu 280	Ser	Туг	. Lys	s Asr	285
Gl	y Ly:	s Pro	g As	9 Gly	y Vai	l Ası	тул	: Ile	295	g Thr	Asp	o Glu	ı Glu	300
As	p Ph	e Ar	g Hi	s Lya	s Se	r Se	r Phe	e Vai	l Ile 310	e 0				